

**Affymetrix gene expression data for the 1771 genes expressed at a PPDE value equal to or greater than 0.99  
For replicate experiments comparing RNA extracted from three normal and three SLS-treated skin samples.**

<b>Affy Probe_ID</b>	<b>Sequence Description</b>
215189_at	Consensus includes gb:X99142.1 /DEF=H.sapiens mRNA for hair keratin, hHb6. /FEA=mRNA /GEN=hHb6 /PROD=type II intermediate filament of hair keratin /DB_XREF=gi:1903219 /UG=Hs.278658 keratin, hair, basic; 6 (monilethrix)
41469_at	Cluster Incl. L10343:Huma elafin gene, complete cds /cds=(516,869) /gb=L10343 /gi=190337 /ug=Hs.L12341 /len=871
204103_at	gb:NM_002984.1 /DEF=Homo sapiens small inducible cytokine A4 (homologous to mouse Mip-1b) (SCYA4), mRNA. /FEA=mRNA /GEN=SCYA4 /PROD=small inducible cytokine A4 (homologous to mouseMip-1b) /DB_XREF=gi:4506844 /UG=Hs.75703 small inducible cytokine A4 (homologous to mouse Mip-1b) /FL=gb:J04130.1 gb:NM_002984.1 gb:M23502.1 gb:M25316.1
219890_at	gb:NM_013252.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5 (CLECSF5), mRNA. /FEA=mRNA /GEN=CLECSF5 /PROD=C-type (calcium dependent,carbohydrate-recognition domain) lectin, superfamilymember 5 /DB_XREF=gi:10281668 /UG=Hs.126355 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5 /FL=gb:NM_013252.1 gb:AF139768.1
203416_at	gb:NM_000560.1 /DEF=Homo sapiens CD53 antigen (CD53), mRNA. /FEA=mRNA /GEN=CD53 /PROD=CD53 antigen /DB_XREF=gi:10834971 /UG=Hs.82212 CD53 antigen /FL=gb:NM_000560.1 gb:M60871.1 gb:M37033.1
201859_at	gb:NM_002727.1 /DEF=Homo sapiens proteoglycan 1, secretory granule (PRG1), mRNA. /FEA=mRNA /GEN=PRG1 /PROD=proteoglycan 1, secretory granule /DB_XREF=gi:4506044 /UG=Hs.1908 proteoglycan 1, secretory granule /FL=gb:J03223.1 gb:NM_002727.1
205828_at	gb:NM_002422.2 /DEF=Homo sapiens matrix metalloproteinase 3 (stromelysin 1, progelatinase) (MMP3), mRNA. /FEA=mRNA /GEN=MMP3 /PROD=matrix metalloproteinase 3 preproprotein /DB_XREF=gi:13027803 /UG=Hs.83326 matrix metalloproteinase 3 (stromelysin 1, progelatinase) /FL=gb:NM_002422.2 gb:J03209.1
216810_at	Consensus includes gb:AJ406939.1 /DEF=Homo sapiens mRNA for keratin associated protein 4.7 (KRTAP4.7 gene). /FEA=mRNA /GEN=KRTAP4.7 /PROD=keratin associated protein 4.7 /DB_XREF=gi:12655451 /UG=Hs.307020 Homo sapiens mRNA for keratin associated protein 4.7 (KRTAP4.7 gene)

Affy Probe_ID	Sequence Description
202878_s_at	gb:NM_012072.2 /DEF=Homo sapiens complement component C1q receptor (C1QR), mRNA. /FEA=mRNA /GEN=C1QR /PROD=complement component C1q receptor /DB_XREF=gi:11496985 /UG=Hs.97199 complement component C1q receptor /FL=gb:NM_012072.2 gb:U94333.1
217897_at	gb:NM_022003.1 /DEF=Homo sapiens FXYD domain-containing ion transport regulator 6 (FXYD6), mRNA. /FEA=mRNA /GEN=FXYD6 /PROD=FXYD domain-containing ion transport regulator6 /DB_XREF=gi:11612654 /UG=Hs.3807 FXYD domain-containing ion transport regulator 6 /FL=gb:NM_022003.1 gb:AL136699.1
203691_at	gb:NM_002638.1 /DEF=Homo sapiens protease inhibitor 3, skin-derived (SKALP) (PI3), mRNA. /FEA=mRNA /GEN=PI3 /PROD=protease inhibitor 3, skin-derived (SKALP) /DB_XREF=gi:4505786 /UG=Hs.112341 protease inhibitor 3, skin-derived (SKALP) /FL=gb:NM_002638.1
207526_s_at	gb:NM_003856.1 /DEF=Homo sapiens interleukin 1 receptor-like 1 (IL1RL1), mRNA. /FEA=mRNA /GEN=IL1RL1 /PROD=interleukin 1 receptor-like 1 /DB_XREF=gi:4507244 /UG=Hs.66 interleukin 1 receptor-like 1 /FL=gb:NM_003856.1
209596_at	gb:AF245505.1 /DEF=Homo sapiens adlican mRNA, complete cds. /FEA=mRNA /PROD=adlican /DB_XREF=gi:9280404 /UG=Hs.72157 DKFZP564I1922 protein /FL=gb:AF245505.1
210873_x_at	gb:U03891.2 /DEF=Homo sapiens phorbolin I mRNA, complete cds. /FEA=mRNA /PROD=phorbolin I /DB_XREF=gi:4895107 /UG=Hs.226307 phorbolin (similar to apolipoprotein B mRNA editing protein) /FL=gb:U03891.2
212646_at	Consensus includes gb:D42043.1 /DEF=Human mRNA for KIAA0084 gene, partial cds. /FEA=mRNA /GEN=KIAA0084 /DB_XREF=gi:577298 /UG=Hs.79123 KIAA0084 protein
215049_x_at	Consensus includes gb:Z22969.1 /DEF=H.sapiens mRNA for M130 antigen cytoplasmic variant 1. /FEA=mRNA /PROD=M130 antigen cytoplasmic variant 1 /DB_XREF=gi:312143 /UG=Hs.74076 CD163 antigen
202957_at	gb:NM_005335.1 /DEF=Homo sapiens hematopoietic cell-specific Lyn substrate 1 (HCLS1), mRNA. /FEA=mRNA /GEN=HCLS1 /PROD=hematopoietic cell-specific Lyn substrate 1 /DB_XREF=gi:4885404 /UG=Hs.14601 hematopoietic cell-specific Lyn substrate 1 /FL=gb:NM_005335.1
203234_at	gb:NM_003364.1 /DEF=Homo sapiens uridine phosphorylase (UP), mRNA. /FEA=mRNA /GEN=UP /PROD=uridine phosphorylase /DB_XREF=gi:4507838 /UG=Hs.77573 uridine phosphorylase /FL=gb:BC001405.1 gb:NM_003364.1

**Affy Probe ID**

**Sequence Description**

219434_at	gb:NM_018643.1 /DEF=Homo sapiens triggering receptor expressed on myeloid cells 1 (TREM1), mRNA. /FEA=mRNA /GEN=TREM1 /PROD=triggering receptor expressed on myeloid cells1 /DB_XREF=gi:8924261 /UG=Hs.283022 triggering receptor expressed on myeloid cells 1 /FL=gb:AF196329.1 gb:NM_018643.1 gb:AF287008.1
204620_s_at	gb:NM_004385.1 /DEF=Homo sapiens chondroitin sulfate proteoglycan 2 (versican) (CSPG2), mRNA. /FEA=mRNA /GEN=CSPG2 /PROD=chondroitin sulfate proteoglycan 2 (versican) /DB_XREF=gi:4758081 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:NM_004385.1
212588_at	Consensus includes gb:Y00062.1 /DEF=Human mRNA for T200 leukocyte common antigen (CD45, LC-A). /FEA=mRNA /DB_XREF=gi:34275 /UG=Hs.170121 protein tyrosine phosphatase, receptor type, C
201785_at	gb:NM_002933.1 /DEF=Homo sapiens ribonuclease, RNase A family, 1 (pancreatic) (RNASE1), mRNA. /FEA=mRNA /GEN=RNASE1 /PROD=ribonuclease, RNase A family, 1 (pancreatic) /DB_XREF=gi:4506546 /UG=Hs.78224 ribonuclease, RNase A family, 1 (pancreatic) /FL=gb:BC005324.1 gb:NM_002933.1 gb:D26129.1
206385_s_at	gb:NM_020987.1 /DEF=Homo sapiens ankyrin 3, node of Ranvier (ankyrin G) (ANK3), transcript variant 1, mRNA. /FEA=mRNA /GEN=ANK3 /PROD=ankyrin 3, isoform 1 /DB_XREF=gi:10947055 /UG=Hs.75893 ankyrin 3, node of Ranvier (ankyrin G) /FL=gb:NM_020987.1 gb:U13616.1
213733_at	Consensus includes gb:BF740152 /FEA=EST /DB_XREF=gi:12066828 /DB_XREF=est:7n13g08.x1 /CLONE=IMAGE:3564495 /UG=Hs.121555 myosin IE
204122_at	gb:NM_003332.1 /DEF=Homo sapiens TYRO protein tyrosine kinase binding protein (TYROBP), mRNA. /FEA=mRNA /GEN=TYROBP /PROD=TYRO protein tyrosine kinase binding protein /DB_XREF=gi:4507754 /UG=Hs.9963 TYRO protein tyrosine kinase binding protein /FL=gb:AF019562.1 gb:NM_003332.1
201721_s_at	gb:NM_006762.1 /DEF=Homo sapiens Lysosomal-associated multispanning membrane protein-5 (LAPTM5), mRNA. /FEA=mRNA /GEN=LAPTM5 /PROD=Lysosomal-associated multispanning membrane protein-5 /DB_XREF=gi:5803055 /UG=Hs.79356 Lysosomal-associated multispanning membrane protein-5 /FL=gb:U51240.1 gb:NM_006762.1 gb:U30498.1
211538_s_at	gb:U56725.1 /DEF=Human heat shock protein mRNA, complete cds. /FEA=mRNA /PROD=heat shock protein /DB_XREF=gi:4204879 /UG=Hs.75452 heat shock 70kD protein 2 /FL=gb:U56725.1
204438_at	gb:NM_002438.1 /DEF=Homo sapiens mannose receptor, C type 1 (MRC1), mRNA. /FEA=mRNA /GEN=MRC1 /PROD=mannose receptor, C type 1 /DB_XREF=gi:4505244 /UG=Hs.75182 mannose receptor, C type 1 /FL=gb:J05550.1 gb:NM_002438.1

**Affy Probe\_ID**

**Sequence Description**

201666\_at gb:NM\_003254.1 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) (TIMP1), mRNA. /FEA=mRNA /GEN=TIMP1 /PROD=tissue inhibitor of metalloproteinase 1 precursor /DB\_XREF=gi:4507508 /UG=Hs.5831 tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) /FL=gb:BC000866.1 gb:M12670.1 gb:M59906.1 gb:NM\_003254.1

205098\_at Consensus includes gb:AI421071 /FEA=EST /DB\_XREF=gi:4267002 /DB\_XREF=est:tf24a02.x1 /CLONE=IMAGE:2097098 /UG=Hs.301921 chemokine (C-C motif) receptor 1 /FL=gb:L09230.1 gb:L10918.1 gb:NM\_001295.1

203637\_s\_at gb:NM\_000381.1 /DEF=Homo sapiens midline 1 (OpitzBBB syndrome) (MID1), mRNA. /FEA=mRNA /GEN=MID1 /PROD=midline 1 /DB\_XREF=gi:4557752 /UG=Hs.27695 midline 1 (OpitzBBB syndrome) /FL=gb:AF269101.1 gb:AF230976.1 gb:AF035360.1 gb:NM\_000381.1

205180\_s\_at gb:NM\_001109.1 /DEF=Homo sapiens a disintegrin and metalloproteinase domain 8 (ADAM8), mRNA. /FEA=mRNA /GEN=ADAM8 /PROD=a disintegrin and metalloproteinase domain 8 precursor /DB\_XREF=gi:4557252 /UG=Hs.86947 a disintegrin and metalloproteinase domain 8 /FL=gb:D26579.1 gb:NM\_001109.1

207850\_at gb:NM\_002090.1 /DEF=Homo sapiens GRO3 oncogene (GRO3), mRNA. /FEA=mRNA /GEN=GRO3 /PROD=GRO3 oncogene /DB\_XREF=gi:4504156 /UG=Hs.89690 GRO3 oncogene /FL=gb:M36821.1 gb:NM\_002090.1

204232\_at gb:NM\_004106.1 /DEF=Homo sapiens Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide (FCER1G), mRNA. /FEA=mRNA /GEN=FCER1G /PROD=Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide precursor /DB\_XREF=gi:4758343 /UG=Hs.743 Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide /FL=gb:M33195.1 gb:NM\_004106.1

217678\_at Consensus includes gb:AA488687 /FEA=EST /DB\_XREF=gi:2216118 /DB\_XREF=est:ab38f03.s1 /CLONE=IMAGE:843101 /UG=Hs.284235 ESTs

217739\_s\_at gb:NM\_005746.1 /DEF=Homo sapiens pre-B-cell colony-enhancing factor (PBEF), mRNA. /FEA=mRNA /GEN=PBEF /PROD=pre-B-cell colony-enhancing factor /DB\_XREF=gi:5031976 /UG=Hs.239138 pre-B-cell colony-enhancing factor /FL=gb:U02020.1 gb:NM\_005746.1

204994\_at gb:NM\_002463.1 /DEF=Homo sapiens myxovirus (influenza) resistance 2, homolog of murine (MX2), mRNA. /FEA=mRNA /GEN=MX2 /PROD=myxovirus (influenza) resistance 2, homolog of murine /DB\_XREF=gi:11342663 /UG=Hs.926 myxovirus (influenza) resistance 2, homolog of murine /FL=gb:NM\_002463.1 gb:M30818.1 gb:M33883.1



Affy Probe_ID	Sequence Description
212915_at	Consensus includes gb:AL569804 /FEA=EST /DB_XREF=gi:12925507 /DB_XREF=est:AL569804 /CLONE=CS0DI002YJ15 (3 prime) /UG=Hs.177635 KIAA1095 protein
206172_at	gb:NM_000640.1 /DEF=Homo sapiens interleukin 13 receptor, alpha 2 (IL13RA2), mRNA. /FEA=mRNA /GEN=IL13RA2 /PROD=interleukin 13 receptor, alpha 2 /DB_XREF=gi:10834991 /UG=Hs.25954 interleukin 13 receptor, alpha 2 /FL=gb:NM_000640.1 gb:U70981.1
214146_s_at	Consensus includes gb:R64130 /FEA=EST /DB_XREF=gi:836009 /DB_XREF=est:yi18h03.s1 /CLONE=IMAGE:139637 /UG=Hs.2164 pro-platelet basic protein (includes platelet basic protein, beta-thromboglobulin, connective tissue-activating peptide III, neutrophil-activating peptide-2)
220088_at	gb:NM_001736.1 /DEF=Homo sapiens complement component 5 receptor 1 (C5a ligand) (C5R1), mRNA. /FEA=mRNA /GEN=C5R1 /PROD=complement component 5 receptor 1 (C5a ligand) /DB_XREF=gi:4502508 /UG=Hs.2161 complement component 5 receptor 1 (C5a ligand) /FL=gb:M62505.1 gb:NM_001736.1
213603_s_at	Consensus includes gb:BE138888 /FEA=EST /DB_XREF=gi:8601388 /DB_XREF=est:xw97c06.x1 /CLONE=IMAGE:2835946 /UG=Hs.301175 HSPC022 protein
206420_at	gb:NM_005849.1 /DEF=Homo sapiens immunoglobulin superfamily, member 6 (IGSF6), mRNA. /FEA=mRNA /GEN=IGSF6 /PROD=immunoglobulin superfamily, member 6 /DB_XREF=gi:5031672 /UG=Hs.135194 immunoglobulin superfamily, member 6 /FL=gb:NM_005849.1
221731_x_at	Consensus includes gb:BF218922 /FEA=EST /DB_XREF=gi:11112418 /DB_XREF=est:601885091F1 /CLONE=IMAGE:4103447 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican)
220978_at	gb:NM_030966.1 /DEF=Homo sapiens keratin associated protein 1.3 (KRTAP1.3), mRNA. /FEA=mRNA /GEN=KRTAP1.3 /PROD=keratin associated protein 1.3 /DB_XREF=gi:13569939 /FL=gb:NM_030966.1
206987_x_at	gb:NM_003862.1 /DEF=Homo sapiens fibroblast growth factor 18 (FGF18), mRNA. /FEA=mRNA /GEN=FGF18 /PROD=fibroblast growth factor 18 /DB_XREF=gi:4503694 /UG=Hs.49585 fibroblast growth factor 18 /FL=gb:AB007422.1 gb:AF075292.1 gb:NM_003862.1
208306_x_at	gb:NM_021983.2 /DEF=Homo sapiens major histocompatibility complex, class II, DR beta 4 (HLA-DRB4), mRNA. /FEA=mRNA /GEN=HLA-DRB4 /PROD=major histocompatibility complex, class II, DRbeta 4 /DB_XREF=gi:11875206 /UG=Hs.293934 major histocompatibility complex, class II, DR beta 4 /FL=gb:NM_021983.2

Affy Probe_ID	Sequence Description
204470_at	gb:NM_001511.1 /DEF=Homo sapiens GRO1 oncogene (melanoma growth stimulating activity, alpha) (GRO1), mRNA. /FEA=mRNA /GEN=GRO1 /PROD=GRO1 oncogene (melanoma growth stimulating activity, alpha) /DB_XREF=gi:4504152 /UG=Hs.789 GRO1 oncogene (melanoma growth stimulating activity, alpha) /FL=gb:NM_001511.1
205419_at	gb:NM_004951.1 /DEF=Homo sapiens Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor) (EBI2), mRNA. /FEA=mRNA /GEN=EBI2 /PROD=Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor) /DB_XREF=gi:4826705 /UG=Hs.784 Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor) /FL=gb:L08177.1 gb:NM_004951.1
220272_at	gb:NM_017637.1 /DEF=Homo sapiens hypothetical protein FLJ20043 (FLJ20043), mRNA. /FEA=mRNA /GEN=FLJ20043 /PROD=hypothetical protein FLJ20043 /DB_XREF=gi:8923050 /UG=Hs.103853 hypothetical protein FLJ20043 /FL=gb:NM_017637.1
203980_at	gb:NM_001442.1 /DEF=Homo sapiens fatty acid binding protein 4, adipocyte (FABP4), mRNA. /FEA=mRNA /GEN=FABP4 /PROD=fatty acid binding protein 4, adipocyte /DB_XREF=gi:4557578 /UG=Hs.83213 fatty acid binding protein 4, adipocyte /FL=gb:BC003672.1 gb:J02874.1 gb:NM_001442.1
218574_s_at	gb:NM_014583.1 /DEF=Homo sapiens LIM and cysteine-rich domains 1 (LMCD1), mRNA. /FEA=mRNA /GEN=LMCD1 /PROD=LIM and cysteine-rich domains 1 /DB_XREF=gi:7657308 /UG=Hs.279943 LIM and cysteine-rich domains 1 /FL=gb:AF216709.1 gb:BC000646.1 gb:AF169284.1 gb:NM_014583.1
210986_s_at	gb:Z24727.1 /DEF=H.sapiens tropomyosin isoform mRNA, complete CDS. /FEA=mRNA /PROD=tropomyosin isoform /DB_XREF=gi:854188 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:Z24727.1
213524_s_at	Consensus includes gb:NM_015714.1 /DEF=Homo sapiens putative lymphocyte G0G1 switch gene (G0S2), mRNA. /FEA=CDS /GEN=G0S2 /PROD=putative lymphocyte G0G1 switch gene /DB_XREF=gi:7657103 /UG=Hs.95910 putative lymphocyte G0G1 switch gene /FL=gb:NM_015714.1
205685_at	Consensus includes gb:BG236280 /FEA=EST /DB_XREF=gi:12750052 /DB_XREF=est:naf24d07.x1 /CLONE=IMAGE:4141884 /UG=Hs.27954 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) /FL=gb:U04343.1 gb:NM_006889.1
201150_s_at	gb:NM_000362.2 /DEF=Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory) (TIMP3), mRNA. /FEA=mRNA /GEN=TIMP3 /PROD=tissue inhibitor of metalloproteinase 3 precursor /DB_XREF=gi:9257248 /UG=Hs.245188 tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory) /FL=gb:U67195.1 gb:U02571.1 gb:U14394.1 gb:NM_000362.2

**Affy Probe ID**

**Sequence Description**

215193_x_at	Consensus includes gb:AJ297586.1 /DEF=Homo sapiens mRNA for MHC class II antigen (HLA-DRB1 gene), DRB1*0402 allele. /FEA=CDS /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:10185079 /UG=Hs.279930 major histocompatibility complex, class II, DR beta 3
217478_s_at	Consensus includes gb:X76775 /DEF=H.sapiens HLA-DMA gene /FEA=mRNA_1 /DB_XREF=gi:512468 /UG=Hs.77522 major histocompatibility complex, class II, DM alpha
202499_s_at	gb:Nm_006931.1 /DEF=Homo sapiens solute carrier family 2 (facilitated glucose transporter), member 3 (SLC2A3), mRNA. /FEA=mRNA /GEN=SLC2A3 /PROD=solute carrier family 2 (facilitated glucosetransporter), member 3 /DB_XREF=gi:5902089 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3 /FL=gb:M20681.1 gb:Nm_006931.1
221698_s_at	gb:AF313468.1 /DEF=Homo sapiens dendritic cell-associated C-type lectin-1 mRNA, complete cds. /FEA=mRNA /PROD=dendritic cell-associated C-type lectin-1 /DB_XREF=gi:13649707 /FL=gb:AF313468.1
210845_s_at	gb:U08839.1 /DEF=Human urokinase-type plasminogen activator receptor mRNA, complete cds. /FEA=mRNA /PROD=urokinase-type plasminogen activator receptor /DB_XREF=gi:517197 /UG=Hs.179657 plasminogen activator, urokinase receptor /FL=gb:U08839.1
204924_at	gb:Nm_003264.1 /DEF=Homo sapiens toll-like receptor 2 (TLR2), mRNA. /FEA=mRNA /GEN=TLR2 /PROD=toll-like receptor2 /DB_XREF=gi:4507528 /UG=Hs.63668 toll-like receptor 2 /FL=gb:U88878.1 gb:AF051152.1 gb:Nm_003264.1
205269_at	Consensus includes gb:AI123251 /FEA=EST /DB_XREF=gi:3539017 /DB_XREF=est:qa47g03.x1 /CLONE=IMAGE:1689940 /UG=Hs.2488 lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD) /FL=gb:Nm_005565.2 gb:U20158.1
201427_s_at	gb:Nm_005410.1 /DEF=Homo sapiens selenoprotein P, plasma, 1 (SEPP1), mRNA. /FEA=mRNA /GEN=SEPP1 /PROD=selenoprotein P precursor /DB_XREF=gi:4885590 /UG=Hs.3314 selenoprotein P, plasma, 1 /FL=gb:Nm_005410.1
219947_at	gb:Nm_016184.1 /DEF=Homo sapiens C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 (CLECSF6), mRNA. /FEA=mRNA /GEN=CLECSF6 /PROD=C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /DB_XREF=gi:7705337 /UG=Hs.115515 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 /FL=gb:AF109146.1 gb:Nm_016184.1 gb:AF067800.1

**Affy Probe ID**

**Sequence Description**

203535_at	gb:NM_002965.2 /DEF=Homo sapiens S100 calcium-binding protein A9 (calgranulin B) (S100A9), mRNA. /FEA=mRNA /GEN=S100A9 /PROD=S100 calcium-binding protein A9 /DB_XREF=gi:9845520 /UG=Hs.112405 S100 calcium-binding protein A9 (calgranulin B) /FL=gb:M26311.1 gb:NM_002965.2
213260_at	Consensus includes gb:AU145890 /FEA=EST /DB_XREF=gi:11007411 /DB_XREF=est:AU145890 /CLONE=HEMBA1006158 /UG=Hs.284186 forkhead box C1 /FL=gb:NM_001453.1
211991_s_at	Consensus includes gb:M27487.1 /DEF=Homo sapiens MHC class II DPw3-alpha-1 chain mRNA, complete cds. /FEA=CDS /GEN=HLA-DPA1 /PROD=MHC class II DP3-alpha /DB_XREF=gi:703088 /UG=Hs.914 Human mRNA for SB classII histocompatibility antigen alpha-chain /FL=gb:M27487.1
204457_s_at	gb:NM_002048.1 /DEF=Homo sapiens growth arrest-specific 1 (GAS1), mRNA. /FEA=mRNA /GEN=GAS1 /PROD=growth arrest-specific 1 /DB_XREF=gi:4503918 /UG=Hs.65029 growth arrest-specific 1 /FL=gb:NM_002048.1 gb:L13698.1
203104_at	gb:NM_005211.1 /DEF=Homo sapiens colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog (CSF1R), mRNA. /FEA=mRNA /GEN=CSF1R /PROD=colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog /DB_XREF=gi:4885158 /UG=Hs.174142 colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog /FL=gb:NM_005211.1
211071_s_at	gb:BC006471.1 /DEF=Homo sapiens, ALL1-fused gene from chromosome 1q, clone MGC:4013, mRNA, complete cds. /FEA=mRNA /PROD=ALL1-fused gene from chromosome 1q /DB_XREF=gi:13623686 /FL=gb:BC006471.1
217764_s_at	gb:AF183421.1 /DEF=Homo sapiens small GTP-binding protein rab22b mRNA, complete cds. /FEA=mRNA /PROD=small GTP-binding protein rab22b /DB_XREF=gi:9963780 /UG=Hs.223025 RAB31, member RAS oncogene family /FL=gb:AF234995.1 gb:BC001148.1 gb:U59877.1 gb:U57091.1 gb:NM_006868.1 gb:AF183421.1
209312_x_at	gb:U65585.1 /DEF=Homo sapiens MHC class II antigen (HLA-DRB1) mRNA, HLA-DRB1*PBL allele, complete cds. /FEA=mRNA /GEN=HLA-DRB1 /PROD=MHC class II antigen /DB_XREF=gi:5478215 /UG=Hs.180255 major histocompatibility complex, class II, DR beta 1 /FL=gb:M33600.1 gb:NM_002124.1 gb:M28583.1 gb:U65585.1
208894_at	gb:M60334.1 /DEF=Human MHC class II HLA-DR-alpha mRNA, complete cds. /FEA=mRNA /GEN=HLA-DRA /PROD=cell surface glycoprotein /DB_XREF=gi:188255 /UG=Hs.76807 major histocompatibility complex, class II, DR alpha /FL=gb:M60334.1 gb:NM_019111.1

Affy Probe_ID	Sequence Description
201422_at	gb:NM_006332.1 /DEF=Homo sapiens interferon, gamma-inducible protein 30 (IFI30), mRNA. /FEA=mRNA /GEN=IFI30 /PROD=interferon, gamma-inducible protein 30 /DB_XREF=gi:5453695 /UG=Hs.14623 interferon, gamma-inducible protein 30 /FL=gb:J03909.1 gb:NM_006332.1 gb:AF097362.1
203932_at	gb:NM_002118.1 /DEF=Homo sapiens major histocompatibility complex, class II, DM beta (HLA-DMB), mRNA. /FEA=mRNA /GEN=HLA-DMB /PROD=major histocompatibility complex, class II, DMbeta /DB_XREF=gi:4504398 /UG=Hs.1162 major histocompatibility complex, class II, DM beta /FL=gb:NM_002118.1 gb:U15085.1
216250_s_at	Consensus includes gb:X77598.1 /DEF=H.sapiens LAM A3 mRNA for laminin alpha 3 chain. /FEA=mRNA /DB_XREF=gi:9716101 /UG=Hs.83450 laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin) gb:NM_002275.1 /DEF=Homo sapiens keratin 15 (KRT15), mRNA. /FEA=mRNA /GEN=KRT15 /PROD=keratin 15 /DB_XREF=gi:4504914 /UG=Hs.80342 keratin 15 /FL=gb:BC002641.1 gb:NM_002275.1
204734_at	gb:U62027.1 /DEF=Human anaphylatoxin C3a receptor (HNFAG09) mRNA, complete cds. /FEA=mRNA /GEN=HNFAG09 /PROD=anaphylatoxin C3a receptor /DB_XREF=gi:1511643 /UG=Hs.155935 complement component 3a receptor 1 /FL=gb:U28488.1 gb:U62027.1 gb:NM_004054.1
201108_s_at	Consensus includes gb:BF05462 /FEA=EST /DB_XREF=gi:10809358 /DB_XREF=est:7j80e06.x1 /CLONE=IMAGE:3392770 /UG=Hs.87409 thrombospondin 1 /FL=gb:NM_003246.1
202054_s_at	gb:NM_000382.1 /DEF=Homo sapiens aldehyde dehydrogenase 3 family, member A2 (ALDH3A2), mRNA. /FEA=mRNA /GEN=ALDH3A2 /PROD=aldehyde dehydrogenase 3A2 /DB_XREF=gi:4557302 /UG=Hs.159608 aldehyde dehydrogenase 3 family, member A2 /FL=gb:L47162.1 gb:U46689.1 gb:NM_000382.1
203936_s_at	gb:NM_004994.1 /DEF=Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA. /FEA=mRNA /GEN=MMP9 /PROD=matrix metalloproteinase 9 preproprotein /DB_XREF=gi:4826835 /UG=Hs.151738 matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) /FL=gb:J05070.1 gb:NM_004994.1
211276_at	gb:AF063606.1 /DEF=Homo sapiens brain my048 protein mRNA, complete cds. /FEA=mRNA /PROD=brain my048 protein /DB_XREF=gi:12002041 /UG=Hs.17481 Homo sapiens clone 24606 mRNA sequence /FL=gb:AF063606.1
202625_at	Consensus includes gb:A1356412 /FEA=EST /DB_XREF=gi:4108033 /DB_XREF=est:qz26h02.x1 /CLONE=IMAGE:2028051 /UG=Hs.80887 v-yes-1 Yamaguchi sarcoma viral related oncogene homolog /FL=gb:NM_002350.1

Affy Probe_ID	Sequence Description
203797_at	gb:AF039555.1 /DEF=Homo sapiens visinin-like protein 1 (VSNL1) mRNA, complete cds. /FEA=mRNA /GEN=VSNL1 /PROD=visinin-like protein 1 /DB_XREF=gi:4104813 /UG=Hs.2288 visinin-like 1 /FL=gb:AF039555.1 gb:NM_003385.1 gb:AB001104.1 gb:U14747.1
206337_at	gb:NM_001838.1 /DEF=Homo sapiens chemokine (C-C motif) receptor 7 (CCR7), mRNA. /FEA=mRNA /GEN=CCR7 /PROD=chemokine (C-C motif) receptor 7 /DB_XREF=gi:4502640 /UG=Hs.1652 chemokine (C-C motif) receptor 7 /FL=gb:L08176.1 gb:NM_001838.1 gb:L31581.1
205686_s_at	gb:NM_006889.1 /DEF=Homo sapiens CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) (CD86), mRNA. /FEA=mRNA /GEN=CD86 /PROD=CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) /DB_XREF=gi:5901919 /UG=Hs.27954 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) /FL=gb:U04343.1 gb:NM_006889.1
221059_s_at	gb:NM_021615.1 /DEF=Homo sapiens carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 (CHST6), mRNA. /FEA=mRNA /GEN=CHST6 /PROD=carbohydrate (N-acetylglucosamine 6-O)sulfotransferase 6 /DB_XREF=gi:11055975 /UG=Hs.157439 carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 6 /FL=gb:AF219990.1 gb:NM_021615.1
220972_s_at	gb:NM_030975.1 /DEF=Homo sapiens keratin associated protein 9.9 (KRTAP9.9), mRNA. /FEA=mRNA /GEN=KRTAP9.9 /PROD=keratin associated protein 9.9 /DB_XREF=gi:13569951 /FL=gb:NM_030975.1
206207_at	gb:NM_001828.3 /DEF=Homo sapiens Charot-Leyden crystal protein (CLC), mRNA. /FEA=mRNA /GEN=CLC /PROD=Charot-Leyden crystal protein /DB_XREF=gi:6325464 /UG=Hs.889 Charot-Leyden crystal protein /FL=gb:L01664.1 gb:NM_001828.3
219093_at	gb:NM_017933.1 /DEF=Homo sapiens hypothetical protein FLJ20701 (FLJ20701), mRNA. /FEA=mRNA /GEN=FLJ20701 /PROD=hypothetical protein FLJ20701 /DB_XREF=gi:8923631 /UG=Hs.169764 hypothetical protein FLJ20701 /FL=gb:NM_017933.1
215223_s_at	Consensus includes gb:W46388 /FEA=EST /DB_XREF=gi:1331076 /DB_XREF=est:zc32c08.s1 /CLONE=IMAGE:324014 /UG=Hs.31885 superoxide dismutase 2, mitochondrial
221491_x_at	Consensus includes gb:AA807056 /FEA=EST /DB_XREF=gi:2876632 /DB_XREF=est:oc35c12.s1 /CLONE=IMAGE:1351702 /UG=Hs.279930 major histocompatibility complex, class II, DR beta 3 /FL=gb:M27635.1 gb:NM_022555.1 gb:M26038.1 gb:U95989.1 gb:U95819.1 gb:U66825.1
204912_at	gb:NM_001558.1 /DEF=Homo sapiens interleukin 10 receptor, alpha (IL10RA), mRNA. /FEA=mRNA /GEN=IL10RA /PROD=interleukin 10 receptor, alpha /DB_XREF=gi:4504632 /UG=Hs.327 interleukin 10 receptor, alpha /FL=gb:NM_001558.1 gb:U00672.1

**Affy Probe ID**

**Sequence Description**

204015_s_at	gb:BC002671.1 /DEF=Homo sapiens, dual specificity phosphatase 4, clone MGC:3713, mRNA, complete cds. /FEA=mRNA /PROD=dual specificity phosphatase 4 /DB_XREF=gi:12803670 /UG=Hs.2359 dual specificity phosphatase 4 /FL=gb:U48807.1 gb:NM_001394.2 gb:BC002671.1 gb:U211108.1
202087_s_at	gb:NM_001912.1 /DEF=Homo sapiens cathepsin L (CTSL), mRNA. /FEA=mRNA /GEN=CTSL /PROD=cathepsin L /DB_XREF=gi:4503154 /UG=Hs.78056 cathepsin L /FL=gb:NM_001912.1
204222_s_at	gb:NM_006851.1 /DEF=Homo sapiens glioma pathogenesis-related protein (RTVP1), mRNA. /FEA=mRNA /GEN=RTVP1 /PROD=glioma pathogenesis-related protein /DB_XREF=gi:5803150 /UG=Hs.64639 glioma pathogenesis-related protein /FL=gb:U16307.1 gb:NM_006851.1
201148_s_at	Consensus includes gb:A W338933 /FEA=EST /DB_XREF=gi:6835559 /DB_XREF=est:ha68g07.x1 /CLONE=IMAGE:2878908 /UG=Hs.245188 tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory) /FL=gb:U67195.1 gb:U02571.1 gb:U14394.1 gb:NM_000362.2
217362_x_at	Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5915893 /UG=Hs.167385 Homo sapiens MHC class II antigen HLA-DRB6 mRNA, partial cds
203305_at	gb:NM_000129.2 /DEF=Homo sapiens coagulation factor XIII, A1 polypeptide (F13A1), mRNA. /FEA=mRNA /GEN=F13A1 /PROD=coagulation factor XIII A1 subunit precursor /DB_XREF=gi:9961355 /UG=Hs.80424 coagulation factor XIII, A1 polypeptide /FL=gb:M14354.1 gb:NM_000129.2
209290_s_at	gb:BC001283.1 /DEF=Homo sapiens, Similar to nuclear factor IB, clone MGC:5146, mRNA, complete cds. /FEA=mRNA /PROD=Similar to nuclear factor IB /DB_XREF=gi:12654878 /UG=Hs.33287 nuclear factor IB /FL=gb:BC001283.1 gb:U85193.1 gb:NM_005596.1
202888_s_at	gb:NM_001150.1 /DEF=Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA. /FEA=mRNA /GEN=ANPEP /PROD=membrane alanine aminopeptidase precursor /DB_XREF=gi:4502094 /UG=Hs.1239 alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) /FL=gb:M22324.1 gb:NM_001150.1
206295_at	gb:NM_001562.1 /DEF=Homo sapiens interleukin 18 (interferon-gamma-inducing factor) (IL18), mRNA. /FEA=mRNA /GEN=IL18 /PROD=interleukin 18 /DB_XREF=gi:4504652 /UG=Hs.83077 interleukin 18 (interferon-gamma-inducing factor) /FL=gb:D49950.1 gb:AF077611.1 gb:NM_001562.1

Affy Probe_ID	Sequence Description
217963_s_at	gb:NM_014380.1 /DEF=Homo sapiens p75NTR-associated cell death executor; ovarian granulosa cell protein (13kD) (DXS6984E), mRNA. /FEA=mRNA /GEN=DXS6984E /PROD=p75NTR-associated cell death executor; ovarian granulosa cell protein (13kD) /DB_XREF=gi:7657043 /UG=Hs.17775 p75NTR-associated cell death executor; ovarian granulosa cell protein (13kD) /FL=gb:NM_014380.1 gb:AF187064.1
205237_at	gb:NM_002003.2 /DEF=Homo sapiens ficolin (collagenfibrinogen domain-containing) 1 (FCN1), mRNA. /FEA=mRNA /GEN=FCN1 /PROD=ficolin 1 precursor /DB_XREF=gi:8051583 /UG=Hs.252136 ficolin (collagenfibrinogen domain-containing) 1 /FL=gb:D83920.1 gb:NM_002003.2
206343_s_at	gb:NM_013959.1 /DEF=Homo sapiens neuregulin 1 (NRG1), transcript variant SMDF, mRNA. /FEA=mRNA /GEN=NRG1 /PROD=neuregulin 1 isoform SMDF /DB_XREF=gi:7669517 /UG=Hs.172816 neuregulin 1 /FL=gb:NM_013959.1 gb:L41827.1
209555_s_at	gb:M98399.1 /DEF=Human antigen CD36 (clone 21) mRNA, complete cds. /FEA=mRNA /GEN=CD36 /PROD=antigen CD36 /DB_XREF=gi:180112 /UG=Hs.75613 CD36 antigen (collagen type I receptor, thrombospondin receptor) /FL=gb:M98399.1
206569_at	gb:NM_006850.1 /DEF=Homo sapiens suppression of tumorigenicity 16 (melanoma differentiation) (ST16), mRNA. /FEA=mRNA /GEN=ST16 /PROD=suppression of tumorigenicity 16 (melanomadifferentiation) /DB_XREF=gi:5803085 /UG=Hs.315463 suppression of tumorigenicity 16 (melanoma differentiation) /FL=gb:U16261.1 gb:NM_006850.1
202341_s_at	Consensus includes gb:AA149745 /FEA=EST /DB_XREF=gi:1720818 /DB_XREF=est:zo02h04.s1 /CLONE=IMAGE:566551 /UG=Hs.12372 tripartite motif protein TRIM2 /FL=gb:AF220018.1 gb:NM_015271.1
205067_at	gb:NM_000576.1 /DEF=Homo sapiens interleukin 1, beta (IL1B), mRNA. /FEA=mRNA /GEN=IL1B /PROD=interleukin 1, beta /DB_XREF=gi:10835144 /UG=Hs.126256 interleukin 1, beta /FL=gb:NM_000576.1 gb:K02770.1 gb:M15330.1 gb:M54933.1
208075_s_at	gb:NM_006273.2 /DEF=Homo sapiens small inducible cytokine A7 (monocyte chemotactic protein 3) (SCYA7), mRNA. /FEA=mRNA /GEN=SCYA7 /PROD=monocyte chemotactic protein 3 precursor /DB_XREF=gi:13435401 /FL=gb:NM_006273.2
219371_s_at	gb:NM_016270.1 /DEF=Homo sapiens Kruppel-like factor (LOC51713), mRNA. /FEA=mRNA /GEN=LOC51713 /PROD=Kruppel-like factor /DB_XREF=gi:7706468 /UG=Hs.107740 Kruppel-like factor 2 (lung) /FL=gb:AF134053.1 gb:AF205849.1 gb:NM_016270.1



**Affy Probe\_ID**

**Sequence Description**

212998\_x\_at Consensus includes gb:AI583173 /FEA=EST /DB\_XREF=gi:4569070 /DB\_XREF=est:tg64e04.x1 /CLONE=IMAGE:2213598 /UG=Hs.73931 major histocompatibility complex, class II, DQ beta 1

205185\_at gb:NM\_006846.1 /DEF=Homo sapiens serine protease inhibitor, Kazal type, 5 (SPINK5), mRNA. /FEA=mRNA /GEN=SPINK5 /PROD=serine protease inhibitor, Kazal type, 5 /DB\_XREF=gi:5803218 /UG=Hs.331555 serine protease inhibitor, Kazal type, 5 /FL=gb:NM\_006846.1

204619\_s\_at Consensus includes gb:BF590263 /FEA=EST /DB\_XREF=gi:11682587 /DB\_XREF=est:nab22b12.x1 /CLONE=IMAGE:3266638 /UG=Hs.81800 chondroitin sulfate proteoglycan 2 (versican) /FL=gb:NM\_004385.1

202901\_x\_at gb:BC002642.1 /DEF=Homo sapiens, cathepsin S, clone MGC:3886, mRNA, complete cds. /FEA=mRNA /PROD=cathepsin S /DB\_XREF=gi:12803614 /UG=Hs.181301 cathepsin S /FL=gb:BC002642.1 gb:M86553.1 gb:NM\_004079.1 gb:M90696.1

204588\_s\_at gb:NM\_003982.1 /DEF=Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 7 (SLC7A7), mRNA. /FEA=mRNA /GEN=SLC7A7 /PROD=solute carrier family 7 (cationic amino acid transporter, y+ system), member 7 /DB\_XREF=gi:4507054 /UG=Hs.194693 solute carrier family 7 (cationic amino acid transporter, y+ system), member 7 /FL=gb:BC003062.1 gb:AF092032.1 gb:NM\_003982.1 gb:AB011263.1

204446\_s\_at gb:NM\_000698.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase (ALOX5), mRNA. /FEA=mRNA /GEN=ALOX5 /PROD=arachidonate 5-lipoxygenase /DB\_XREF=gi:4502056 /UG=Hs.89499 arachidonate 5-lipoxygenase /FL=gb:J03600.1 gb:J03571.1 gb:NM\_000698.1

204670\_x\_at gb:NM\_002125.1 /DEF=Homo sapiens major histocompatibility complex, class II, DR beta 5 (HLA-DRB5), mRNA. /FEA=mRNA /GEN=HLA-DRB5 /PROD=major histocompatibility complex, class II, DR beta 5 /DB\_XREF=gi:4504412 /UG=Hs.308026 major histocompatibility complex, class II, DR beta 5 /FL=gb:M11867.1 gb:M20430.1 gb:NM\_002125.1

209606\_at gb:L06633.1 /DEF=Human transcription factor mRNA, complete cds. /FEA=mRNA /PROD=transcription factor /DB\_XREF=gi:431327 /UG=Hs.270 plectstrin homology, Sec7 and coiledcoil domains, binding protein /FL=gb:AF068836.1 gb:L06633.1 gb:NM\_004288.2

214467\_at Consensus includes gb:NM\_003608.1 /DEF=Homo sapiens G protein-coupled receptor 65 (GPR65), mRNA. /FEA=CDS /GEN=GPR65 /PROD=G protein-coupled receptor 65 /DB\_XREF=gi:4507420 /UG=Hs.131924 G protein-coupled receptor 65 /FL=gb:NM\_003608.1

211742\_s\_at gb:BC005926.1 /DEF=Homo sapiens, ecotropic viral integration site 2B, clone MGC:14529, mRNA, complete cds. /FEA=mRNA /PROD=ecotropic viral integration site 2B /DB\_XREF=gi:13543535 /FL=gb:BC005926.1

Affy Probe_ID	Sequence Description
221477_s_at	Consensus includes gb:BF575213 /FEA=EST /DB_XREF=gi:11648925 /DB_XREF=est:602133624F1 /CLONE=IMAGE:4288756 /UG=Hs.177781 Homo sapiens, clone MGC:5618, mRNA, complete cds /FL=gb:BC001980.1
203961_at	Consensus includes gb:AL157398 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the nebulin protein (NEBL, actin-binding Z-disc protein) /FEA=mRNA_1 /DB_XREF=gi:10045326 /UG=Hs.5025 nebulin /FL=gb:NM_006393.1
204735_at	gb:NM_006202.1 /DEF=Homo sapiens phosphodiesterase 4A, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E2) (PDE4A), mRNA. /FEA=mRNA /GEN=PDE4A /PROD=phosphodiesterase 4A, cAMP-specific (dunce(Drosophila)-homolog phosphodiesterase E2) /DB_XREF=gi:5453861 /UG=Hs.89901 phosphodiesterase 4A, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E2) /FL=gb:U97584.1 gb:U68532.1 gb:NM_006202.1
204057_at	Consensus includes gb:AI073984 /FEA=EST /DB_XREF=gi:3400628 /DB_XREF=est:oy66c05.x1 /CLONE=IMAGE:1670792 /UG=Hs.14453 interferon consensus sequence binding protein 1 /FL=gb:M91196.1 gb:NM_002163.1
214036_at	Consensus includes gb:BE464799 /FEA=EST /DB_XREF=gi:9510493 /DB_XREF=est:hs87a06.x1 /CLONE=IMAGE:3144178 /UG=Hs.27342 ESTs
204661_at	gb:NM_001803.1 /DEF=Homo sapiens CDW52 antigen (CAMPATH-1 antigen) (CDW52), mRNA. /FEA=mRNA /GEN=CDW52 /PROD=CDW52 antigen (CAMPATH-1 antigen) /DB_XREF=gi:4502760 /UG=Hs.276770 CDW52 antigen (CAMPATH-1 antigen) /FL=gb:BC000644.1 gb:NM_001803.1
204992_s_at	gb:NM_002628.1 /DEF=Homo sapiens profilin 2 (PFN2), mRNA. /FEA=mRNA /GEN=PFN2 /PROD=profilin 2 /DB_XREF=gi:4505750 /UG=Hs.91747 profilin 2 /FL=gb:L10678.1 gb:NM_002628.1
201278_at	Consensus includes gb:N21202 /FEA=EST /DB_XREF=gi:1126372 /DB_XREF=est:yx41e12.s1 /CLONE=IMAGE:264334 /UG=Hs.81988 disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein) /FL=gb:U39050.1 gb:U53446.1 gb:BC003064.1 gb:NM_001343.1
210895_s_at	gb:L25259.1 /DEF=Human CTLA4 counter-receptor (B7-2) mRNA, complete cds. /FEA=mRNA /GEN=B7-2 /PROD=CTLA4 counter-receptor /DB_XREF=gi:416368 /UG=Hs.27954 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) /FL=gb:L25259.1
202910_s_at	gb:NM_001784.1 /DEF=Homo sapiens CD97 antigen (CD97), mRNA. /FEA=mRNA /GEN=CD97 /PROD=CD97 antigen /DB_XREF=gi:4502690 /UG=Hs.3107 CD97 antigen /FL=gb:NM_001784.1

**Affy Probe\_ID**

**Sequence Description**

216834_at	Consensus includes gb:S59049.1 /DEF=BL34=B cell activation gene human, mRNA, 1398 nt. /FEA=mRNA /GEN=BL34 /DB_XREF=gi:299704 /UG=Hs.75256 regulator of G-protein signalling 1
207067_s_at	gb:Nm_002112.1 /DEF=Homo sapiens histidine decarboxylase (HDC), mRNA. /FEA=mRNA /GEN=HDC /PROD=histidine decarboxylase /DB_XREF=gi:4504364 /UG=Hs.1481 histidine decarboxylase /FL=gb:M60445.1 gb:Nm_002112.1
208981_at	Consensus includes gb:AA702701 /FEA=EST /DB_XREF=gi:2705814 /DB_XREF=est:zi90h02.s1 /CLONE=IMAGE:448083 /UG=Hs.78146 plateletendothelial cell adhesion molecule (CD31 antigen) /FL=gb:M37780.1 gb:M28526.1 gb:Nm_000442.1
204173_at	gb:Nm_002475.1 /DEF=Homo sapiens myosin, light polypeptide 1, alkali; skeletal, fast (MYL1), mRNA. /FEA=mRNA /GEN=MYL1 /PROD=myosin, light polypeptide 1, alkali; skeletal, fast /DB_XREF=gi:4505302 /UG=Hs.90318 myosin, light polypeptide 1, alkali; skeletal, fast /FL=gb:M31211.1 gb:Nm_002475.1
213029_at	Consensus includes gb:BG478428 /FEA=EST /DB_XREF=gi:13410807 /DB_XREF=est:602523839F1 /CLONE=IMAGE:4642353 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKFZp564H1916 (from clone DKFZp564H1916)
209921_at	gb:AB040875.1 /DEF=Homo sapiens hxCT mRNA for cystineglutamate exchanger, complete cds. /FEA=mRNA /GEN=hxCT /PROD=cystineglutamate exchanger /DB_XREF=gi:13516845 /UG=Hs.6682 solute carrier family 7, (cationic amino acid transporter, y+ system) member 11 /FL=gb:AB040875.1
203474_at	gb:Nm_006633.1 /DEF=Homo sapiens IQ motif containing GTPase activating protein 2 (IQGAP2), mRNA. /FEA=mRNA /GEN=IQGAP2 /PROD=IQ motif containing GTPase activating protein 2 /DB_XREF=gi:5729886 /UG=Hs.78993 IQ motif containing GTPase activating protein 2 /FL=gb:U51903.1 gb:Nm_006633.1
205372_at	gb:Nm_002655.1 /DEF=Homo sapiens pleiomorphic adenoma gene 1 (PLAG1), mRNA. /FEA=mRNA /GEN=PLAG1 /PROD=pleiomorphic adenoma gene 1 /DB_XREF=gi:4505854 /UG=Hs.14968 pleiomorphic adenoma gene 1 /FL=gb:U65002.1 gb:Nm_002655.1
211429_s_at	gb:AF119873.1 /DEF=Homo sapiens PRO2275 mRNA, complete cds. /FEA=mRNA /PROD=PRO2275 /DB_XREF=gi:7770182 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /FL=gb:AF119873.1
216841_s_at	Consensus includes gb:X15132.1 /DEF=Human mRNA for manganese containing superoxide dismutase (EC 1.15.1.1). /FEA=mRNA /DB_XREF=gi:34794 /UG=Hs.318885 superoxide dismutase 2, mitochondrial

**Affy Probe\_ID**

**Sequence Description**

219142\_at gb:NM\_023940.1 /DEF=Homo sapiens hypothetical protein MGC2827 (MGC2827), mRNA. /FEA=mRNA /GEN=MGC2827 /PROD=hypothetical protein MGC2827 /DB\_XREF=gi:13027611 /UG=Hs.8035 hypothetical protein MGC2827 /FL=gb:BC001087.1 gb:BC001846.1 gb:NM\_023940.1

209723\_at gb:BC002538.1 /DEF=Homo sapiens, serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9, clone MGC:2131, mRNA, complete cds. /FEA=mRNA /PROD=serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 /DB\_XREF=gi:12803428 /UG=Hs.104879 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 /FL=gb:L40378.1 gb:BC002538.1 gb:U71364.1 gb:NM\_004155.1

204475\_at gb:NM\_002421.2 /DEF=Homo sapiens matrix metalloproteinase 1 (interstitial collagenase) (MMP1), mRNA. /FEA=mRNA /GEN=MMP1 /PROD=matrix metalloproteinase 1 preproprotein /DB\_XREF=gi:13027798 /UG=Hs.83169 matrix metalloproteinase 1 (interstitial collagenase) /FL=gb:NM\_002421.2 gb:M13509.1

205987\_at gb:NM\_001765.1 /DEF=Homo sapiens CD1C antigen, c polypeptide (CD1C), mRNA. /FEA=mRNA /GEN=CD1C /PROD=CD1C antigen, c polypeptide /DB\_XREF=gi:4502646 /UG=Hs.1311 CD1C antigen, c polypeptide /FL=gb:M28827.1 gb:NM\_001765.1

206488\_s\_at gb:NM\_000072.1 /DEF=Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor) (CD36), mRNA. /FEA=mRNA /GEN=CD36 /PROD=CD36 antigen (collagen type I receptor, thrombospondin receptor) /DB\_XREF=gi:4557418 /UG=Hs.75613 CD36 antigen (collagen type I receptor, thrombospondin receptor) /FL=gb:M24795.1 gb:M98398.1 gb:L06850.1 gb:NM\_000072.1

203153\_at gb:NM\_001548.1 /DEF=Homo sapiens interferon-induced protein with tetratricopeptide repeats 1 (IFIT1), mRNA. /FEA=mRNA /GEN=IFIT1 /PROD=interferon-induced protein with tetratricopeptide repeats 1 /DB\_XREF=gi:4504584 /UG=Hs.20315 interferon-induced protein with tetratricopeptide repeats 1 /FL=gb:M24594.1 gb:NM\_001548.1

205003\_at gb:NM\_014705.1 /DEF=Homo sapiens KIAA0716 gene product (KIAA0716), mRNA. /FEA=mRNA /GEN=KIAA0716 /PROD=KIAA0716 gene product /DB\_XREF=gi:7662263 /UG=Hs.118140 KIAA0716 gene product /FL=gb:AB018259.1 gb:NM\_014705.1

204440\_at gb:NM\_004233.1 /DEF=Homo sapiens CD83 antigen (activated B lymphocytes, immunoglobulin superfamily) (CD83), mRNA. /FEA=mRNA /GEN=CD83 /PROD=CD83 antigen (activated B lymphocytes, immunoglobulin superfamily) /DB\_XREF=gi:4757945 /UG=Hs.79197 CD83 antigen (activated B lymphocytes, immunoglobulin superfamily) /FL=gb:NM\_004233.1

**Affy Probe ID**

**Sequence Description**

213988\_s\_at Consensus includes gb:BE971383 /FEA=EST /DB\_XREF=gi:10584719 /DB\_XREF=est:601651561F1 /CLONE=IMAGE:3934935 /UG=Hs.28491 spermidinespermine N1-acetyltransferase

205986\_at gb:NM\_004920.1 /DEF=Homo sapiens apoptosis-associated tyrosine kinase (AATK), mRNA. /FEA=mRNA /GEN=AATK /PROD=apoptosis-associated tyrosine kinase /DB\_XREF=gi:4758641 /UG=Hs.128316 apoptosis-associated tyrosine kinase /FL=gb:AB014541.1 gb:NM\_004920.1

212992\_at Consensus includes gb:AI935123 /FEA=EST /DB\_XREF=gi:5673993 /DB\_XREF=est:wp13h09.x1 /CLONE=IMAGE:2464769 /UG=Hs.57548 ESTs

211002\_s\_at gb:AF230389.1 /DEF=Homo sapiens tripartite motif protein TRIM29 beta mRNA, complete cds. /FEA=mRNA /PROD=tripartite motif protein TRIM29 beta /DB\_XREF=gi:12275865 /UG=Hs.82237 ataxia-telangiectasia group D-associated protein /FL=gb:AF230389.1

213924\_at Consensus includes gb:BF476502 /FEA=EST /DB\_XREF=gi:11547329 /DB\_XREF=est:naa27a03.x1 /CLONE=IMAGE:3255844 /UG=Hs.154145 hypothetical protein FLJ11585

209619\_at gb:K01144.1 /DEF=Human major histocompatibility class II antigen gamma chain mRNA, complete cds. /FEA=mRNA /PROD=class II antigen gamma chain /DB\_XREF=gi:188469 /UG=Hs.84298 CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) /FL=gb:NM\_004355.1 gb:K01144.1

200648\_s\_at gb:NM\_002065.1 /DEF=Homo sapiens glutamate-ammonia ligase (glutamine synthase) (GLUL), mRNA. /FEA=mRNA /GEN=GLUL /PROD=glutamate-ammonia ligase (glutamine synthase) /DB\_XREF=gi:4504026 /UG=Hs.170171 glutamate-ammonia ligase (glutamine synthase) /FL=gb:NM\_002065.1

203038\_at gb:NM\_002844.1 /DEF=Homo sapiens protein tyrosine phosphatase, receptor type, K (PTPRK), mRNA. /FEA=mRNA /GEN=PTPRK /PROD=protein tyrosine phosphatase, receptor type, K /DB\_XREF=gi:4506316 /UG=Hs.79005 protein tyrosine phosphatase, receptor type, K /FL=gb:L77886.1 gb:NM\_002844.1

207710\_at gb:NM\_014357.1 /DEF=Homo sapiens skin-specific protein (XP5), mRNA. /FEA=mRNA /GEN=XP5 /PROD=skin-specific protein /DB\_XREF=gi:7657684 /UG=Hs.234766 skin-specific protein /FL=gb:AF005080.1 gb:NM\_014357.1

205400\_at gb:NM\_000377.1 /DEF=Homo sapiens Wiskott-Aldrich syndrome (eczema-thrombocytopenia) (WAS), mRNA. /FEA=mRNA /GEN=WAS /PROD=Wiskott-Aldrich syndrome protein /DB\_XREF=gi:4507908 /UG=Hs.2157 Wiskott-Aldrich syndrome (eczema-thrombocytopenia) /FL=gb:NM\_000377.1 gb:U12707.1 gb:U19927.1

222088\_s\_at Consensus includes gb:AA778684 /FEA=EST /DB\_XREF=gi:2838015 /DB\_XREF=est:af87g07.s1 /CLONE=1049052 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3

Affy Probe ID	Sequence Description
205863_at	gb:NM_005621.1 /DEF=Homo sapiens S100 calcium-binding protein A12 (calgranulin C) (S100A12), mRNA. /FEA=mRNA /GEN=S100A12 /PROD=S100 calcium-binding protein A12 /DB_XREF=gi:5032058 /UG=Hs.19413 S100 calcium-binding protein A12 (calgranulin C) /FL=gb:D83664.1 gb:D49549.1 gb:NM_005621.1
210020_x_at	gb:M58026.1 /DEF=Human NB-1 mRNA, complete cds. /FEA=mRNA /GEN=NB-1 /DB_XREF=gi:189080 /UG=Hs.239600 calmodulin-like 3 /FL=gb:M36707.1 gb:M58026.1 gb:NM_005185.1
204724_s_at	gb:NM_001853.1 /DEF=Homo sapiens collagen, type IX, alpha 3 (COL9A3), mRNA. /FEA=mRNA /GEN=COL9A3 /PROD=collagen, type IX, alpha 3 /DB_XREF=gi:4502966 /UG=Hs.53563 collagen, type IX, alpha 3 /FL=gb:L41162.1 gb:NM_001853.1
220016_at	gb:NM_024060.1 /DEF=Homo sapiens hypothetical protein MGC5395 (MGC5395), mRNA. /FEA=mRNA /GEN=MGC5395 /PROD=hypothetical protein MGC5395 /DB_XREF=gi:13129033 /UG=Hs.165215 hypothetical protein MGC5395 /FL=gb:BC000926.1 gb:NM_024060.1
212154_at	Consensus includes gb:AI380298 /FEA=EST /DB_XREF=gi:4190151 /DB_XREF=est:tf99h06.x1 /CLONE=IMAGE:2107451 /UG=Hs.1501 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
217762_s_at	Consensus includes gb:BE789881 /FEA=EST /DB_XREF=gi:10211079 /DB_XREF=est:601476750F1 /CLONE=IMAGE:3879493 /UG=Hs.223025 RAB31, member RAS oncogene family /FL=gb:AF234995.1 gb:BC001148.1 gb:U59877.1 gb:U57091.1 gb:NM_006868.1 gb:AF183421.1
206214_at	gb:NM_005084.1 /DEF=Homo sapiens phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma) (PLA2G7), mRNA. /FEA=mRNA /GEN=PLA2G7 /PROD=phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma) /DB_XREF=gi:4826883 /UG=Hs.93304 phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma) /FL=gb:U24577.1 gb:NM_005084.1 gb:U20157.1
202037_s_at	gb:NM_003012.2 /DEF=Homo sapiens secreted frizzled-related protein 1 (SFRP1), mRNA. /FEA=mRNA /GEN=SFRP1 /PROD=secreted frizzled-related protein 1 /DB_XREF=gi:8400731 /UG=Hs.7306 secreted frizzled-related protein 1 /FL=gb:AF001900.1 gb:AF017987.1 gb:AF056087.1 gb:NM_003012.2
209289_at	Consensus includes gb:AI700518 /FEA=EST /DB_XREF=gi:4988418 /DB_XREF=est:we37d09.x1 /CLONE=IMAGE:2343281 /UG=Hs.33287 nuclear factor IB /FL=gb:BC001283.1 gb:U85193.1 gb:NM_005596.1

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## Sequence Description

209949\_at gb:BC001606.1 /DEF=Homo sapiens, Similar to neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2), clone MGC:2275, mRNA, complete cds. /FEA=mRNA /PROD=Similar to neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2) /DB\_XREF=gi:12804408 /UG=Hs.949 neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2) /FL=gb:BC001606.1 gb:M32011.1 gb:NM\_000433.1 gb:NM\_000418.1 /DEF=Homo sapiens interleukin 4 receptor (IL4R), mRNA. /FEA=mRNA /GEN=IL4R /PROD=interleukin 4 receptor precursor /DB\_XREF=gi:4557668 /UG=Hs.75545 interleukin 4 receptor /FL=gb:NM\_000418.1

201426\_s\_at Consensus includes gb:AI922599 /FEA=EST /DB\_XREF=gi:5658563 /DB\_XREF=est:wm90b11.x1 /CLONE=IMAGE:2443197 /UG=Hs.297753 vimentin /FL=gb:BC000163.2 gb:NM\_003380.1

203561\_at gb:NM\_021642.1 /DEF=Homo sapiens Fc fragment of IgG, low affinity IIa, receptor for (CD32) (FCGR2A), mRNA. /FEA=mRNA /GEN=FCGR2A /PROD=Fc fragment of IgG, low affinity IIa, receptor for (CD32) /DB\_XREF=gi:11056051 /UG=Hs.78864 Fc fragment of IgG, low affinity IIa, receptor for (CD32) /FL=gb:NM\_021642.1 gb:M31932.1 gb:J03619.1 gb:M28697.1

201675\_at gb:NM\_003488.1 /DEF=Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), mRNA. /FEA=mRNA /GEN=AKAP1 /PROD=A kinase (PRKA) anchor protein 1 /DB\_XREF=gi:4502014 /UG=Hs.78921 A kinase (PRKA) anchor protein 1 /FL=gb:BC000729.1 gb:NM\_003488.1

217989\_at gb:NM\_016245.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR2 (LOC511170), mRNA. /FEA=mRNA /GEN=LOC511170 /PROD=retinal short-chain dehydrogenasereductaseretSDR2 /DB\_XREF=gi:7705904 /UG=Hs.12150 retinal short-chain dehydrogenasereductase retSDR2 /FL=gb:AF126780.1 gb:NM\_016245.1

219403\_s\_at gb:NM\_006665.1 /DEF=Homo sapiens heparanase (HPSE), mRNA. /FEA=mRNA /GEN=HPSE /PROD=heparanase /DB\_XREF=gi:5729872 /UG=Hs.44227 heparanase /FL=gb:AF165154.1 gb:AF152376.1 gb:NM\_006665.1 gb:AF084467.1 gb:AF155510.1

214382\_at Consensus includes gb:AL021331 /DEF=Human DNA sequence from clone 366N23 on chromosome 6q27. Contains two genes similar to consecutive parts of the C. elegans UNC-93 (protein 1, C46F11.1) gene, a KIAA0173 and Tubulin-Tyrosine Ligase LIKE gene, a Mitotic Feedback Control Protein MADP... /FEA=mRNA\_2 /DB\_XREF=gi:3355532 /UG=Hs.22033 Human DNA sequence from clone 366N23 on chromosome 6q27. Contains two genes similar to consecutive parts of the C. elegans UNC-93 (protein 1, C46F11.1) gene, a KIAA0173 and Tubulin-Tyrosine Ligase LIKE gene, a Mitotic Feedback Control Protein MADP2 Homolo

# Affy Probe\_ID

## Sequence Description

210987\_x\_at gb:M19267.1 /DEF=Human tropomyosin mRNA, complete cds. /FEA=mRNA /DB\_XREF=gi:339943 /UG=Hs.77899 tropomyosin 1 (alpha) /FL=gb:M19267.1  
209685\_s\_at gb:M13975.1 /DEF=Homo sapiens protein kinase C beta-II type (PRKCB1) mRNA, complete cds. /FEA=mRNA /GEN=PRKCB1 /PROD=protein kinase C beta-II type /DB\_XREF=gi:189968 /UG=Hs.77202 protein kinase C, beta 1 /FL=gb:M13975.1  
208018\_s\_at gb:NM\_002110.1 /DEF=Homo sapiens hemopoietic cell kinase (HCK); mRNA. /FEA=mRNA /GEN=HCK /PROD=hemopoietic cell kinase /DB\_XREF=gi:4504356 /UG=Hs.89555 hemopoietic cell kinase /FL=gb:M16591.1 gb:NM\_002110.1  
201667\_at gb:NM\_000165.2 /DEF=Homo sapiens gap junction protein, alpha 1, 43kD (connexin 43) (GJA1), mRNA. /FEA=mRNA /GEN=GJA1 /PROD=connexin 43 /DB\_XREF=gi:4755136 /UG=Hs.74471 gap junction protein, alpha 1, 43kD (connexin 43) /FL=gb:M65188.1 gb:NM\_000165.2  
211361\_s\_at Consensus includes gb:AJ001696.2 /DEF=Homo sapiens mRNA for hurpin, clone R7-1.1. /FEA=mRNA /GEN=PII3 /PROD=hurpin /DB\_XREF=gi:6018505 /UG=Hs.241407 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13 /FL=gb:AF169949.1 gb:NM\_012397.1  
204949\_at gb:NM\_002162.2 /DEF=Homo sapiens intercellular adhesion molecule 3 (ICAM3), mRNA. /FEA=mRNA /GEN=ICAM3 /PROD=intercellular adhesion molecule 3 precursor /DB\_XREF=gi:12545399 /UG=Hs.99995 intercellular adhesion molecule 3 /FL=gb:NM\_002162.2  
212975\_at Consensus includes gb:AB020677.2 /DEF=Homo sapiens mRNA for KIAA0870 protein, partial cds. /FEA=mRNA /GEN=KIAA0870 /PROD=KIAA0870 protein /DB\_XREF=gi:6635136 /UG=Hs.18166 KIAA0870 protein  
201842\_s\_at Consensus includes gb:AI826799 /FEA=EST /DB\_XREF=gi:5447470 /DB\_XREF=est:wk56d07.xl /CLONE=IMAGE:2419405 /UG=Hs.76224 EGF-containing fibulin-like extracellular matrix protein 1 /FL=gb:U03877.1 gb:NM\_004105.2  
210644\_s\_at gb:AF109683.1 /DEF=Homo sapiens leukocyte-associated Ig-like receptor 1b mRNA, complete cds. /FEA=mRNA /PROD=leukocyte-associated Ig-like receptor 1b /DB\_XREF=gi:6563041 /UG=Hs.115808 leukocyte-associated Ig-like receptor 1 /FL=gb:AF251509.2 gb:NM\_021706.1 gb:AF109683.1  
218002\_s\_at gb:NM\_004887.1 /DEF=Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BRAK) (SCYB14), mRNA. /FEA=mRNA /GEN=SCYB14 /PROD=small inducible cytokine subfamily B(Cys-X-Cys), member 14 (BRAK) /DB\_XREF=gi:4757869 /UG=Hs.24395 small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BRAK) /FL=gb:BC003513.1 gb:AF073957.1 gb:NM\_004887.1 gb:AF144103.1 gb:AF106911.1



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**Sequence Description**

205270_s_at	gb:NM_005565.2 /DEF=Homo sapiens lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD) (LCP2), mRNA. /FEA=mRNA /GEN=LCP2 /PROD=lymphocyte cytosolic protein 2 /DB_XREF=gi:7382491 /UG=Hs.2488 lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD) /FL=gb:NM_005565.2 gb:U20158.1
222242_s_at	Consensus includes gb:AF243527 /DEF=Homo sapiens serine protease gene cluster, complete sequence /FEA=CDS_12 /DB_XREF=gi:11244757 /UG=Hs.50915 kallikrein 5
210757_x_at	gb:AF188298.1 /DEF=Homo sapiens disabled 2 p93 (DAB2) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=DAB2 /PROD=disabled 2 p93 /DB_XREF=gi:6176335 /UG=Hs.81988 disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein) /FL=gb:AF188298.1
203903_s_at	gb:NM_014799.1 /DEF=Homo sapiens hephaestin (HEPH), mRNA. /FEA=mRNA /GEN=HEPH /PROD=hephaestin /DB_XREF=gi:7662253 /UG=Hs.31720 hephaestin /FL=gb:AB014598.1 gb:NM_014799.1
201790_s_at	Consensus includes gb:AW150953 /FEA=EST /DB_XREF=gi:6198851 /DB_XREF=est:yg42f09.x1 /CLONE=IMAGE:2630249 /UG=Hs.11806 7-dehydrocholesterol reductase /FL=gb:BC000054.1 gb:AF034544.1 gb:AF067127.1 gb:AF096305.1 gb:NM_001360.1
209879_at	Consensus includes gb:AI741056 /FEA=EST /DB_XREF=gi:5109344 /DB_XREF=est:wg25h08.x1 /CLONE=IMAGE:2366175 /UG=Hs.79283 selectin P ligand /FL=gb:U02297.1 gb:NM_003006.2
201910_at	Consensus includes gb:BF213279 /FEA=EST /DB_XREF=gi:11106865 /DB_XREF=est:601844779F1 /CLONE=IMAGE:4070203 /UG=Hs.183738 FERM, RhoGEF (ARHGEF) and plectstrin domain protein 1 (chondrocyte-derived) /FL=gb:AB008430.1 gb:NM_005766.1
215704_at	Consensus includes gb:AL356504 /DEF=Human DNA sequence from clone RP1-14N1 on chromosome 1q21.1-21.3 Contains ESTs, GSSs and STSs. Contains the FLG gene for profilaggrin and part of a gene for a novel S-100CaBP type calcium binding domain protein similar to trichohyalin.n /FEA=mRNA /DB_XREF=gi:9588237 /UG=Hs.73995 filaggrin
201858_s_at	gb:J03223.1 /DEF=Human secretory granule proteoglycan peptide core mRNA, complete cds. /FEA=mRNA /GEN=PRG1 /DB_XREF=gi:190419 /UG=Hs.1908 proteoglycan 1, secretory granule /FL=gb:J03223.1 gb:NM_002727.1
210095_s_at	gb:M31159.1 /DEF=Human growth hormone-dependent insulin-like growth factor-binding protein mRNA, complete cds. /FEA=mRNA /GEN=IGFBP1 /DB_XREF=gi:183115 /UG=Hs.77326 insulin-like growth factor binding protein 3 /FL=gb:BC000013.1 gb:M31159.1

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## Sequence Description

203320\_at gb:NM\_005475.1 /DEF=Homo sapiens lymphocyte adaptor protein (LNK), mRNA. /FEA=mRNA /GEN=LNK /PROD=lymphocyte adaptor protein /DB\_XREF=gi:4885454 /UG=Hs.13131 lymphocyte adaptor protein /FL=gb:AF055581.1 gb:NM\_005475.1

201487\_at gb:NM\_001814.1 /DEF=Homo sapiens cathepsin C (CTSC), mRNA. /FEA=mRNA /GEN=CTSC /PROD=cathepsin C /DB\_XREF=gi:4503140 /UG=Hs.10029 cathepsin C /FL=gb:NM\_001814.1

205147\_x\_at gb:NM\_000631.2 /DEF=Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 1, mRNA. /FEA=mRNA /GEN=NCF4 /PROD=neutrophil cytosolic factor 4 (40kD), isoform 1 /DB\_XREF=gi:7382494 /UG=Hs.196352 neutrophil cytosolic factor 4 (40kD) /FL=gb:AB025220.1 gb:NM\_000631.2

202672\_s\_at gb:NM\_001674.1 /DEF=Homo sapiens activating transcription factor 3 (ATF3), mRNA. /FEA=mRNA /GEN=ATF3 /PROD=activating transcription factor 3 long isoform /DB\_XREF=gi:4502262 /UG=Hs.460 activating transcription factor 3 /FL=gb:L19871.1 gb:NM\_001674.1

201791\_s\_at gb:NM\_001360.1 /DEF=Homo sapiens 7-dehydrocholesterol reductase (DHCR7), mRNA. /FEA=mRNA /GEN=DHCR7 /PROD=7-dehydrocholesterol reductase /DB\_XREF=gi:4503320 /UG=Hs.11806 7-dehydrocholesterol reductase /FL=gb:BC000054.1 gb:AF034544.1 gb:AF067127.1 gb:AF096305.1 gb:NM\_001360.1

202856\_s\_at gb:NM\_004207.1 /DEF=Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 3 (SLC16A3), mRNA. /FEA=mRNA /GEN=SLC16A3 /PROD=solute carrier family 16 (monocarboxylic acid transporters), member 3 /DB\_XREF=gi:4759111 /UG=Hs.85838 solute carrier family 16 (monocarboxylic acid transporters), member 3 /FL=gb:U81800.1 gb:NM\_004207.1

213587\_s\_at Consensus includes gb:AI884867 /FEA=EST /DB\_XREF=gi:5590031 /DB\_XREF=est:wl85g03.x1 /CLONE=IMAGE:2431732 /UG=Hs.91379 ribosomal protein L26

214598\_at Consensus includes gb:AL049977.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C122 (from clone DKFZp564C122). /FEA=mRNA /DB\_XREF=gi:4884227 /UG=Hs.162209 claudin 8 /FL=gb:NM\_012132.1

218959\_at gb:NM\_017409.1 /DEF=Homo sapiens homeo box C10 (HOXC10), mRNA. /FEA=mRNA /GEN=HOXC10 /PROD=homeo box C10 /DB\_XREF=gi:8393550 /UG=Hs.44276 homeo box C10 /FL=gb:BC001293.1 gb:AF255675.1 gb:NM\_017409.1

202391\_at gb:NM\_006317.1 /DEF=Homo sapiens brain acid-soluble protein 1 (BASP1), mRNA. /FEA=mRNA /GEN=BASP1 /PROD=brain acid-soluble protein 1 /DB\_XREF=gi:5453749 /UG=Hs.79516 brain abundant, membrane attached signal protein 1 /FL=gb:BC000518.1 gb:AF039656.1 gb:NM\_006317.1

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## Sequence Description

219607\_s\_at gb:NM\_024021.1 /DEF=Homo sapiens membrane-spanning 4-domains, subfamily A, member 4 (MS4A4), mRNA. /FEA=mRNA /GEN=MS4A4 /PROD=membrane-spanning 4-domains, subfamily A, member4 /DB\_XREF=gi:13430865 /UG=Hs.325960 membrane-spanning 4-domains, subfamily A, member 4 /FL=gb:AB013102.1 gb:NM\_024021.1 gb:AF068288.1 gb:NM\_016650.1

202086\_at gb:NM\_002462.1 /DEF=Homo sapiens myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78) (MX1), mRNA. /FEA=mRNA /GEN=MX1 /PROD=myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78) /DB\_XREF=gi:4505290 /UG=Hs.76391 myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78) /FL=gb:M30817.1 gb:M33882.1 gb:NM\_002462.1

213032\_at Consensus includes gb:AI186739 /FEA=EST /DB\_XREF=gi:3737377 /DB\_XREF=est:qe79c01.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKFZp564H1916 (from clone DKFZp564H1916)

210785\_s\_at gb:AB035482.1 /DEF=Homo sapiens mRNA for ICB-1beta, complete cds. /FEA=mRNA /PROD=ICB-1beta /DB\_XREF=gi:8272423 /UG=Hs.10649 basement membrane-induced gene /FL=gb:AB035482.1

220413\_at gb:NM\_014579.1 /DEF=Homo sapiens zinc transporter (ZIP2), mRNA. /FEA=mRNA /GEN=ZIP2 /PROD=zinc transporter /DB\_XREF=gi:7657698 /UG=Hs.175783 zinc transporter /FL=gb:AF186081.1 gb:D83039.1 gb:NM\_014579.1

202158\_s\_at gb:NM\_006561.1 /DEF=Homo sapiens CUG triplet repeat, RNA-binding protein 2 (CUGBP2), mRNA. /FEA=mRNA /GEN=CUGBP2 /PROD=CUG triplet repeat, RNA-binding protein 2 /DB\_XREF=gi:5729815 /UG=Hs.211610 CUG triplet repeat, RNA-binding protein 2 /FL=gb:U69546.1 gb:AF036956.1 gb:AF090694.1 gb:NM\_006561.1

218736\_s\_at gb:NM\_017734.1 /DEF=Homo sapiens hypothetical protein FLJ20271 (FLJ20271), mRNA. /FEA=mRNA /GEN=FLJ20271 /PROD=hypothetical protein FLJ20271 /DB\_XREF=gi:8923242 /UG=Hs.14606 hypothetical protein FLJ20271 /FL=gb:NM\_017734.1

219247\_s\_at gb:NM\_024630.1 /DEF=Homo sapiens hypothetical protein FLJ20984 (FLJ20984), mRNA. /FEA=mRNA /GEN=FLJ20984 /PROD=hypothetical protein FLJ20984 /DB\_XREF=gi:13375857 /UG=Hs.38270 hypothetical protein FLJ20984 /FL=gb:NM\_024630.1

206380\_s\_at gb:NM\_002621.1 /DEF=Homo sapiens properdin P factor, complement (PFC), mRNA. /FEA=mRNA /GEN=PFC /PROD=properdin P factor, complement /DB\_XREF=gi:4505736 /UG=Hs.53155 properdin P factor, complement /FL=gb:NM\_002621.1 gb:M83652.1

Affy Probe_ID	Sequence Description
208438_s_at	gb:NM_005248.1 /DEF=Homo sapiens Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog (FGR), mRNA. /FEA=mRNA /GEN=FGR /PROD=Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog /DB_XREF=gi:4885234 /UG=Hs.1422 Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog /FL=gb:M19722.1 gb:NM_005248.1
213241_at	Consensus includes gb:AF035307.1 /DEF=Homo sapiens clone 23785 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661068 /UG=Hs.184697 Homo sapiens clone 23785 mRNA sequence
210065_s_at	gb:AB002155.1 /DEF=Homo sapiens mRNA for human uroplakin Ib, complete cds. /FEA=mRNA /PROD=human uroplakin Ib /DB_XREF=gi:6682736 /UG=Hs.271580 uroplakin 1B /FL=gb:AB015234.1 gb:AF042331.1 gb:NM_006952.1 gb:AB002155.1
214866_at	Consensus includes gb:X74039.1 /DEF=H.sapiens mRNA for urokinase plasminogen activator receptor. /FEA=mRNA /PROD=urokinase plasminogen activator receptor /DB_XREF=gi:456192 /UG=Hs.179657 plasminogen activator, urokinase receptor
204436_at	gb:NM_025201.1 /DEF=Homo sapiens hypothetical protein PP1628 (PP1628), mRNA. /FEA=mRNA /GEN=PP1628 /PROD=hypothetical protein PP1628 /DB_XREF=gi:13376795 /UG=Hs.7159 hypothetical protein PP1628 /FL=gb:NM_025201.1
214438_at	Consensus includes gb:M60721.1 /DEF=Human homeobox gene, complete cds. /FEA=mRNA /DB_XREF=gi:183789 /UG=Hs.74870 H2.0 (Drosophila)-like homeo box 1 /FL=gb:NM_021958.1 gb:M60721.1
219511_s_at	gb:NM_005460.1 /DEF=Homo sapiens synuclein, alpha interacting protein (synphilin) (SNCAIP), mRNA. /FEA=mRNA /GEN=SNCAIP /PROD=synuclein alpha interacting protein /DB_XREF=gi:4885602 /UG=Hs.24948 synuclein, alpha interacting protein (synphilin) /FL=gb:AF076929.1 gb:NM_005460.1
209969_s_at	gb:BC002704.1 /DEF=Homo sapiens, Similar to signal transducer and activator of transcription 1, 91kD, clone MGC:3493, mRNA, complete cds. /FEA=mRNA /PROD=Similar to signal transducer and activator of transcription 1, 91kD /DB_XREF=gi:12803734 /UG=Hs.21486 signal transducer and activator of transcription 1, 91kD /FL=gb:BC002704.1
218854_at	gb:NM_013352.1 /DEF=Homo sapiens squamous cell carcinoma antigen recognized by T cell (SART-2), mRNA. /FEA=mRNA /GEN=SART-2 /PROD=squamous cell carcinoma antigen recognized by T cell /DB_XREF=gi:7019520 /UG=Hs.58636 squamous cell carcinoma antigen recognized by T cell /FL=gb:AF098066.1 gb:NM_013352.1

Affy Probe_ID	Sequence Description
204858_s_at	gb:NM_001953.2 /DEF=Homo sapiens endothelial cell growth factor 1 (platelet-derived) (ECGF1), mRNA. /FEA=mRNA /GEN=ECGF1 /PROD=endothelial cell growth factor 1(platelet-derived) /DB_XREF=gi:7669488 /UG=Hs.73946 endothelial cell growth factor 1 (platelet-derived) /FL=gb:NM_001953.2
211936_at	Consensus includes gb:AF216292.1 /DEF=Homo sapiens endoplasmic reticulum luminal Ca2+ binding protein grp78 mRNA, complete cds. /FEA=CDS /PROD=endoplasmic reticulum luminal Ca2+ binding protein grp78 /DB_XREF=gi:7229461 /UG=Hs.75410 heat shock 70kD protein 5 (glucose-regulated protein, 78kD) /FL=gb:AF216292.1
211756_at	gb:BC005961.1 /DEF=Homo sapiens, parathyroid hormone-like hormone, clone MGC:14611, mRNA, complete cds. /FEA=mRNA /PROD=parathyroid hormone-like hormone /DB_XREF=gi:13543620 /FL=gb:BC005961.1
218532_s_at	gb:NM_019000.1 /DEF=Homo sapiens hypothetical protein (FLJ20152), mRNA. /FEA=mRNA /GEN=FLJ20152 /PROD=hypothetical protein /DB_XREF=gi:9506660 /UG=Hs.82273 hypothetical protein /FL=gb:NM_019000.1
203339_at	Consensus includes gb:AI887457 /FEA=EST /DB_XREF=gi:5592621 /DB_XREF=est:wm05f07.x1 /CLONE=IMAGE:2435077 /UG=Hs.179866 solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /FL=gb:NM_003705.1
211133_x_at	gb:AF009643.1 /DEF=Homo sapiens clone 6 immunoglobulin-like transcript 5 protein mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 5 protein /DB_XREF=gi:2662445 /UG=Hs.105928 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 /FL=gb:AF009643.1
217967_s_at	gb:AF288391.1 /DEF=Homo sapiens C1orf24 mRNA, complete cds. /FEA=mRNA /PROD=C1orf24 /DB_XREF=gi:12620191 /UG=Hs.48778 niban protein /FL=gb:AB050477.1 gb:NM_022083.1 gb:AF288391.1
207571_x_at	gb:NM_004848.1 /DEF=Homo sapiens basement membrane-induced gene (ICB-1), mRNA. /FEA=mRNA /GEN=ICB-1 /PROD=basement membrane-induced gene /DB_XREF=gi:4758579 /UG=Hs.10649 basement membrane-induced gene /FL=gb:AF044896.1 gb:NM_004848.1
204493_at	gb:NM_001196.1 /DEF=Homo sapiens BH3 interacting domain death agonist (BID), mRNA. /FEA=mRNA /GEN=BID /PROD=BH3 interacting domain death agonist /DB_XREF=gi:4557360 /UG=Hs.172894 BH3 interacting domain death agonist /FL=gb:AF042083.1 gb:NM_001196.1
214247_s_at	Consensus includes gb:AU148057 /FEA=EST /DB_XREF=gi:11009578 /DB_XREF=est:AU148057 /CLONE=MAMMA1002489 /UG=Hs.278503 regulated in glioma

**Affy Probe ID**

**Sequence Description**

204580\_at gb:NM\_002426.1 /DEF=Homo sapiens matrix metalloproteinase 12 (macrophage elastase) (MMP12), mRNA. /FEA=mRNA /GEN=MMP12 /PROD=matrix metalloproteinase 12 preproprotein /DB\_XREF=gi:4505206 /UG=Hs.1695 matrix metalloproteinase 12 (macrophage elastase) /FL=gb:L23808.1 gb:NM\_002426.1

218009\_s\_at gb:NM\_003981.1 /DEF=Homo sapiens protein regulator of cytokinesis 1 (PRC1), mRNA. /FEA=mRNA /GEN=PRC1 /PROD=protein regulator of cytokinesis 1 /DB\_XREF=gi:4506038 /UG=Hs.5101 protein regulator of cytokinesis 1 /FL=gb:BC003138.1 gb:AF044588.1 gb:NM\_003981.1

203645\_s\_at gb:NM\_004244.1 /DEF=Homo sapiens CD163 antigen (CD163), mRNA. /FEA=mRNA /GEN=CD163 /PROD=CD163 antigen /DB\_XREF=gi:4758721 /UG=Hs.74076 CD163 antigen /FL=gb:NM\_004244.1

207339\_s\_at gb:NM\_002341.1 /DEF=Homo sapiens lymphotoxin beta (TNF superfamily, member 3) (LTB), transcript variant 1, mRNA. /FEA=mRNA /GEN=LTB /PROD=lymphotoxin-beta isoform a /DB\_XREF=gi:4505034 /UG=Hs.890 lymphotoxin beta (TNF superfamily, member 3) /FL=gb:L11015.1 gb:NM\_002341.1

204464\_s\_at gb:NM\_001957.1 /DEF=Homo sapiens endothelin receptor type A (EDNRA), mRNA. /FEA=mRNA /GEN=EDNRA /PROD=endothelin receptor type A /DB\_XREF=gi:4503464 /UG=Hs.76252 endothelin receptor type A /FL=gb:L06622.1 gb:NM\_001957.1

205466\_s\_at gb:NM\_005114.1 /DEF=Homo sapiens heparan sulfate (glucosamine) 3-O-sulfotransferase 1 (HS3ST1), mRNA. /FEA=mRNA /GEN=HS3ST1 /PROD=heparan sulfate D-glucosaminyl3-O-sulfotransferase 1 precursor /DB\_XREF=gi:4826763 /UG=Hs.40968 heparan sulfate (glucosamine) 3-O-sulfotransferase 1 /FL=gb:AF019386.1 gb:NM\_005114.1

222218\_s\_at Consensus includes gb:AJ400843.1 /DEF=Homo sapiens partial mRNA for immunoglobulin-like cell surface receptor FDF03-M14, soluble alternative form. /FEA=mRNA /GEN=FDF03-M14 /PROD=cell surface receptor FDF03-M14 /DB\_XREF=gi:9715838 /UG=Hs.122591 paired immunoglobulin-like receptor alpha

206707\_x\_at gb:NM\_015864.1 /DEF=Homo sapiens chromosome 6 open reading frame 32 (C6ORF32), mRNA. /FEA=mRNA /GEN=C6ORF32 /PROD=PL48 /DB\_XREF=gi:7705341 /UG=Hs.101359 chromosome 6 open reading frame 32 /FL=gb:BC001232.1 gb:U49187.1 gb:NM\_015864.1

200635\_s\_at Consensus includes gb:AU145351 /FEA=EST /DB\_XREF=gi:11006872 /DB\_XREF=est:AU145351 /CLONE=HEMBA1004591 /UG=Hs.75216 protein tyrosine phosphatase, receptor type, F /FL=gb:NM\_002840.1

211434\_s\_at gb:AF015524.1 /DEF=Homo sapiens putative chemokine receptor (GRAM-A) mRNA, complete cds. /FEA=mRNA /GEN=GRAM-A /PROD=putative chemokine receptor /DB\_XREF=gi:3550066 /UG=Hs.302043 chemokine (C-C motif) receptor-like 2 /FL=gb:AF015524.1

# Affy Probe\_ID

## Sequence Description

216565_x_at	Consensus includes gb:AL121994 /DEF=Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U)), STSs and GSSs /FEA=CDS /DB_XREF=gi:8648917 /UG=Hs.302117 Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon induced transmembrane protein 3 (1-8U)), STSs and GSSs
204174_at	gb:NM_001629.1 /DEF=Homo sapiens arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA. /FEA=mRNA /GEN=ALOX5AP /PROD=arachidonate 5-lipoxygenase-activating protein /DB_XREF=gi:4502058 /UG=Hs.100194 arachidonate 5-lipoxygenase-activating protein /FL=gb:NM_001629.1
205483_s_at	gb:NM_005101.1 /DEF=Homo sapiens interferon-stimulated protein, 15 kDa (ISG15), mRNA. /FEA=mRNA /GEN=ISG15 /PROD=interferon-stimulated protein, 15 kDa /DB_XREF=gi:4826773 /UG=Hs.833 interferon-stimulated protein, 15 kDa /FL=gb:M13755.1 gb:NM_005101.1
213036_x_at	Consensus includes gb:Y15724 /DEF=Homo sapiens SERCA3 gene, exons 1-7 (and joined CDS) /FEA=mRNA /DB_XREF=gi:3021395 /UG=Hs.5541 ATPase, Ca++ transporting, ubiquitous
201417_at	Consensus includes gb:AL136179 /DEF=Human DNA sequence from clone RP3-322L4 on chromosome 6. Contains the SOX4 gene for SRY (sex determining region Y)-box 4, a pseudogene similar to predicted fly, worm and yeast genes, ESTs, STSs, GSSs and four CpG islands /FEA=mRNA /DB_XREF=gi:8649149 /UG=Hs.83484 SRY (sex determining region Y)-box 4 /FL=gb:NM_003107.1
210946_at	gb:AF014403.1 /DEF=Homo sapiens type-2 phosphatidic acid phosphatase alpha-2 (PAP2-a2) mRNA, complete cds. /FEA=mRNA /GEN=PAP2-a2 /PROD=type-2 phosphatidic acid phosphatase alpha-2 /DB_XREF=gi:3123849 /UG=Hs.41569 phosphatidic acid phosphatase type 2A /FL=gb:AF014403.1
202053_s_at	gb:L47162.1 /DEF=Human fatty aldehyde dehydrogenase (FALDH) mRNA, complete cds. /FEA=mRNA /GEN=FALDH /PROD=fatty aldehyde dehydrogenase /DB_XREF=gi:1082035 /UG=Hs.159608 aldehyde dehydrogenase 3 family, member A2 /FL=gb:L47162.1 gb:U46689.1 gb:NM_000382.1
205114_s_at	gb:NM_002983.1 /DEF=Homo sapiens small inducible cytokine A3 (homologous to mouse Mip-1a) (SCYA3), mRNA. /FEA=mRNA /GEN=SCYA3 /PROD=small inducible cytokine A3 (homologous to mouse Mip-1a) /DB_XREF=gi:4506842 /UG=Hs.73817 small inducible cytokine A3 (homologous to mouse Mip-1a) /FL=gb:M23452.1 gb:D00044.1 gb:NM_002983.1 gb:M25315.1

Affy Probe_ID	Sequence Description
209173_at	gb:AF088867.1 /DEF=Homo sapiens putative secreted protein XAG mRNA, complete cds. /FEA=mRNA /PROD=putative secreted protein XAG /DB_XREF=gi:6652811 /UG=Hs.91011 anterior gradient 2 (Xenopus laevis) homolog /FL=gb:AF007791.1 gb:AF038451.1 gb:NM_006408.1 gb:AF088867.1
204400_at	gb:NM_005864.1 /DEF=Homo sapiens signal transduction protein (SH3 containing) (EFS2), mRNA. /FEA=mRNA /GEN=EFS2 /PROD=signal transduction protein (SH3 containing) /DB_XREF=gi:5031680 /UG=Hs.24587 signal transduction protein (SH3 containing) /FL=gb:AB001466.1 gb:NM_005864.1
203700_s_at	gb:NM_013989.1 /DEF=Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 1, mRNA. /FEA=mRNA /GEN=DIO2 /PROD=deiodinase, iodothyronine, type II /DB_XREF=gi:7549802 /UG=Hs.154424 deiodinase, iodothyronine, type II /FL=gb:U53506.1 gb:AF093774.1 gb:NM_013989.1
203922_s_at	Consensus includes gb:AI308863 /FEA=EST /DB_XREF=gi:4003734 /DB_XREF=est:qo78d02.x1 /CLONE=IMAGE:1914627 /UG=Hs.88974 cytochrome b-245, beta polypeptide (chronic granulomatous disease) /FL=gb:NM_000397.2
210724_at	gb:AF239764.1 /DEF=Homo sapiens EGF-like module-containing mucin-like receptor EMR3 mRNA, complete cds. /FEA=mRNA /PROD=EGF-like module-containing mucin-like receptor EMR3 /DB_XREF=gi:13183148 /UG=Hs.326777 Homo sapiens EGF-like module-containing mucin-like receptor EMR3 mRNA, complete cds /FL=gb:AF239764.1
214085_x_at	Consensus includes gb:AI912583 /FEA=EST /DB_XREF=gi:5632438 /DB_XREF=est:we11d05.x1 /CLONE=IMAGE:2340777 /UG=Hs.154762 HIV-1 rev binding protein 2
202289_s_at	gb:NM_006997.1 /DEF=Homo sapiens transforming, acidic coiled-coil containing protein 2 (TACC2), mRNA. /FEA=mRNA /GEN=TACC2 /PROD=transforming, acidic coiled-coil containing protein 2 /DB_XREF=gi:11119413 /UG=Hs.272023 transforming, acidic coiled-coil containing protein 2 /FL=gb:AF095791.2 gb:NM_006997.1
213380_x_at	Consensus includes gb:AA911235 /FEA=EST /DB_XREF=gi:3050525 /DB_XREF=est:ol49d02.s1 /CLONE=IMAGE:1526787 /UG=Hs.278657 macrophage stimulating 1 (hepatocyte growth factor-like)
205159_at	Consensus includes gb:AV756141 /FEA=EST /DB_XREF=gi:10913989 /DB_XREF=est:AV756141 /CLONE=BMFAKF10 /UG=Hs.285401 colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage) /FL=gb:NM_000395.1 gb:M59941.1
215966_x_at	Consensus includes gb:AA292874 /FEA=EST /DB_XREF=gi:1941855 /DB_XREF=est:zt66a07.r1 /CLONE=IMAGE:727284 /UG=Hs.1466 glycerol kinase



**Affy Probe ID**

**Sequence Description**

219282_s_at	gb:NM_015930.1 /DEF=Homo sapiens vanilloid receptor-like protein 1 (VRL-1), mRNA. /FEA=mRNA /GEN=VRL-1 /PROD=vanilloid receptor-like protein 1 /DB_XREF=gi:7706764 /UG=Hs.279746 vanilloid receptor-like protein 1 /FL=gb:AF129112.1 gb:NM_015930.1
206932_at	gb:NM_003956.1 /DEF=Homo sapiens cholesterol 25-hydroxylase (CH25H), mRNA. /FEA=mRNA /GEN=CH25H /PROD=cholesterol 25-hydroxylase /DB_XREF=gi:4502498 /UG=Hs.194687 cholesterol 25-hydroxylase /FL=gb:AF059214.1 gb:NM_003956.1
201195_s_at	gb:AB018009.1 /DEF=Homo sapiens mRNA for L-type amino acid transporter 1, complete cds. /FEA=mRNA /GEN=hLAT1 /PROD=L-type amino acid transporter 1 /DB_XREF=gi:5926731 /UG=Hs.184601 solute carrier family 7 (cationic amino acid transporter, y+ system), member 5 /FL=gb:AF077866.1 gb:AB018542.1 gb:AF104032.1 gb:NM_003486.1 gb:AB017908.1 gb:AB018009.1
39402_at	Cluster Incl. M15330:Human interleukin 1-beta (IL1B) mRNA, complete cds /cds=(86,895) /gb=M15330 /gi=186283 /ug=Hs.126256 /len=1497
205786_s_at	gb:NM_000632.2 /DEF=Homo sapiens integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide) (ITGAM), mRNA. /FEA=mRNA /GEN=ITGAM /PROD=integrin alpha M precursor /DB_XREF=gi:6006013 /UG=Hs.172631 integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide) /FL=gb:J03925.1 gb:NM_000632.2
48031_r_at	Cluster Incl. H93077:yv05a04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-241806 /clone_end=3 /gb=H93077 /gi=1099405 /ug=Hs.10235 /len=604
215679_at	Consensus includes gb:AL109706.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362430. /FEA=mRNA /DB_XREF=gi:5689823 /UG=Hs.32769 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362430
204014_at	gb:NM_001394.2 /DEF=Homo sapiens dual specificity phosphatase 4 (DUSP4), mRNA. /FEA=mRNA /GEN=DUSP4 /PROD=dual specificity phosphatase 4 /DB_XREF=gi:12707552 /UG=Hs.2359 dual specificity phosphatase 4 /FL=gb:U48807.1 gb:NM_001394.2 gb:BC002671.1 gb:U21108.1
203761_at	gb:NM_006748.1 /DEF=Homo sapiens Src-like-adaptor (SLA), mRNA. /FEA=mRNA /GEN=SLA /PROD=Src-like-adaptor /DB_XREF=gi:5803170 /UG=Hs.75367 Src-like-adaptor /FL=gb:U30473.1 gb:D89077.1 gb:U44403.1 gb:NM_006748.1

Affy Probe_ID	Sequence Description
201125_s_at	gb:NM_002213.1 /DEF=Homo sapiens integrin, beta 5 (ITGB5), mRNA. /FEA=mRNA /GEN=ITGB5 /PROD=integrin, beta 5 /DB_XREF=gi:4504772 /UG=Hs.149846 integrin, beta 5 /FL=gb:M35011.1 gb:J05633.1 gb:NM_002213.1
208146_s_at	gb:NM_031311.1 /DEF=Homo sapiens serine carboxypeptidase vitellogenic-like (LOC54504), mRNA. /FEA=mRNA /GEN=LOC54504 /PROD=serine carboxypeptidase vitellogenic-like /DB_XREF=gi:13786124 /FL=gb:NM_031311.1
203756_at	gb:NM_014786.1 /DEF=Homo sapiens KIAA0337 gene product (KIAA0337), mRNA. /FEA=mRNA /GEN=KIAA0337 /PROD=KIAA0337 gene product /DB_XREF=gi:7662063 /UG=Hs.45180 KIAA0337 gene product /FL=gb:AB002335.1 gb:NM_014786.1
217028_at	Consensus includes gb:AJ224869 /DEF=Homo sapiens CXCR4 gene encoding receptor CXCR4 /FEA=mRNA /DB_XREF=gi:3059119 /UG=Hs.89414 chemokine (C-X-C motif), receptor 4 (fusin)
217272_s_at	Consensus includes gb:AJ001698.1 /DEF=Homo sapiens mRNA for hurpin, clone R7-16.1. /FEA=mRNA /GEN=PI13 /PROD=hurpin /DB_XREF=gi:6018509 /UG=Hs.241407 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13
218975_at	gb:NM_015719.1 /DEF=Homo sapiens collagen, type V, alpha 3 (COL5A3), mRNA. /FEA=mRNA /GEN=COL5A3 /PROD=procollagen, type V, alpha 3 /DB_XREF=gi:7656988 /UG=Hs.235368 collagen, type V, alpha 3 /FL=gb:AF177941.1 gb:NM_015719.1
219572_at	gb:NM_017954.1 /DEF=Homo sapiens hypothetical protein FLJ20761 (FLJ20761), mRNA. /FEA=mRNA /GEN=FLJ20761 /PROD=hypothetical protein FLJ20761 /DB_XREF=gi:8923671 /UG=Hs.107872 hypothetical protein FLJ20761 /FL=gb:NM_017954.1
207720_at	gb:NM_000427.1 /DEF=Homo sapiens loridin (LOR), mRNA. /FEA=mRNA /GEN=LOR /PROD=loridin /DB_XREF=gi:4557724 /UG=Hs.251680 loridin /FL=gb:M61120.1 gb:NM_000427.1
219666_at	gb:NM_022349.1 /DEF=Homo sapiens CD20-like precursor (LOC64166), mRNA. /FEA=mRNA /GEN=LOC64166 /PROD=CD20-like precursor /DB_XREF=gi:11641258 /UG=Hs.17914 membrane-spanning 4-domains, subfamily A, member6 /FL=gb:AF142409.1 gb:NM_022349.1
201279_s_at	gb:BC003064.1 /DEF=Homo sapiens, disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein), clone MGC:1764, mRNA, complete cds. /FEA=mRNA /PROD=disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein) /DB_XREF=gi:13111753 /UG=Hs.81988 disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein) /FL=gb:U39050.1 gb:U53446.1 gb:BC003064.1 gb:NM_001343.1

**Affy Probe ID**

**Sequence Description**

204007_at	gb:J04162.1 /DEF=Human leukocyte IgG receptor (Fc-gamma-R) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:183036 /UG=Hs.176663 Fc fragment of IgG, low affinity IIb, receptor for (CD16) /FL=gb:NM_000570.1 gb:J04162.1 gb:M24854.1 gb:AB025256.1
212727_at	Consensus includes gb:AB033058.1 /DEF=Homo sapiens mRNA for KIAA1232 protein, partial cds. /FEA=mRNA /GEN=KIAA1232 /PROD=KIAA1232 protein /DB_XREF=gi:6330721 /UG=Hs.11101 KIAA1232 protein
200712_s_at	Consensus includes gb:AI633566 /FEA=EST /DB_XREF=gi:4684896 /DB_XREF=est:th68f09.x1 /CLONE=IMAGE:2123849 /UG=Hs.234279 microtubule-associated protein, RPEB family, member 1 /FL=gb:NM_012325.1 gb:U24166.1
206969_at	gb:NM_021013.2 /DEF=Homo sapiens keratin, hair, acidic, 4 (KRTHA4), mRNA. /FEA=mRNA /GEN=KRTHA4 /PROD=type I hair keratin 4 /DB_XREF=gi:11641406 /UG=Hs.296942 keratin, hair, acidic, 4 /FL=gb:NM_021013.2
219561_at	gb:NM_016429.1 /DEF=Homo sapiens COP2 for nonclathrin coat protein zeta-COP (LOC51226), mRNA. /FEA=mRNA /GEN=LOC51226 /PROD=COP2 for nonclathrin coat protein zeta-COP /DB_XREF=gi:7705982 /UG=Hs.37482 COP2 for nonclathrin coat protein zeta-COP /FL=gb:AB047849.1 gb:AB037938.1 gb:NM_016429.1
201843_s_at	gb:NM_004105.2 /DEF=Homo sapiens EGF-containing fibulin-like extracellular matrix protein 1 (EFEMP1), transcript variant 1, mRNA. /FEA=mRNA /GEN=EFEMP1 /PROD=EGF-containing fibulin-like extracellular matrix protein 1 precursor, isoform a precursor /DB_XREF=gi:9665261 /UG=Hs.76224 EGF-containing fibulin-like extracellular matrix protein 1 /FL=gb:U03877.1 gb:NM_004105.2
203881_s_at	gb:NM_004010.1 /DEF=Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 (DMD), transcript variant Dp427p2, mRNA. /FEA=mRNA /GEN=DMD /PROD=dystrophin Dp427p2 isoform /DB_XREF=gi:5032314 /UG=Hs.169470 dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272 /FL=gb:NM_004010.1
209774_x_at	gb:M57731.1 /DEF=Human gro-beta mRNA, complete cds. /FEA=mRNA /GEN=gro-beta /PROD=cytokine gro-beta /DB_XREF=gi:183626 /UG=Hs.75765 GRO2 oncogene /FL=gb:M57731.1 gb:M36820.1 gb:NM_002089.1
204774_at	gb:NM_014210.1 /DEF=Homo sapiens ecotropic viral integration site 2A (EVI2A), mRNA. /FEA=mRNA /GEN=EVI2A /PROD=ecotropic viral integration site 2A /DB_XREF=gi:7657074 /UG=Hs.70499 ecotropic viral integration site 2A /FL=gb:NM_014210.1

Affy Probe_ID	Sequence Description
203910_at	gb:NM_004815.1 /DEF=Homo sapiens PTPL1-associated RhoGAP 1 (PARG1), mRNA. /FEA=mRNA /GEN=PARG1 /PROD=PTPL1-associated RhoGAP 1 /DB_XREF=gi:4758881 /UG=Hs:70983 PTPL1-associated RhoGAP 1 /FL=gb:U90920.1 gb:NM_004815.1
215726_s_at	Consensus includes gb:M22976.1 /DEF=Human cytochrome b5 mRNA, 3 end. /FEA=mRNA /GEN=CYB5 /PROD=cytochrome b5 /DB_XREF=gi:181228 /UG=Hs:83834 cytochrome b-5
207677_s_at	gb:NM_013416.1 /DEF=Homo sapiens neutrophil cytosolic factor 4 (40kD) (NCF4), transcript variant 2, mRNA. /FEA=mRNA /GEN=NCF4 /PROD=neutrophil cytosolic factor 4 (40kD), isoform 2 /DB_XREF=gi:7382492 /UG=Hs:196352 neutrophil cytosolic factor 4 (40kD) /FL=gb:BC002798.1 gb:AB025219.1 gb:NM_013416.1
220412_x_at	gb:NM_005714.1 /DEF=Homo sapiens potassium channel, subfamily K, member 7 (KCNK7), mRNA. /FEA=mRNA /GEN=KCNK7 /PROD=potassium channel, subfamily K, member 7 /DB_XREF=gi:5031820 /UG=Hs:175218 potassium channel, subfamily K, member 7 /FL=gb:AF110524.1 gb:NM_005714.1
218477_at	gb:NM_014051.1 /DEF=Homo sapiens PTD011 protein (PTD011), mRNA. /FEA=mRNA /GEN=PTD011 /PROD=PTD011 protein /DB_XREF=gi:7662638 /UG=Hs:94896 PTD011 protein /FL=gb:AF078864.1 gb:AF239771.1 gb:NM_014051.1
202643_s_at	Consensus includes gb:AI738896 /FEA=EST /DB_XREF=gi:5100877 /DB_XREF=est:wi22g02.x1 /CLONE=IMAGE:2391026 /UG=Hs:211600 tumor necrosis factor, alpha-induced protein 3 /FL=gb:M59465.1 gb:NM_006290.1
212365_at	Consensus includes gb:BF215996 /FEA=EST /DB_XREF=gi:11109582 /DB_XREF=est:601881549F1 /CLONE=IMAGE:4093740 /UG=Hs:121576 Homo sapiens cDNA FLJ20153 fis, clone COL08656, highly similar to AJ001381 Homo sapiens incomplete cDNA for a mutated allele
204183_s_at	Consensus includes gb:AI478542 /FEA=EST /DB_XREF=gi:4371768 /DB_XREF=est:tm53c11.x1 /CLONE=IMAGE:2161844 /UG=Hs:13944 adrenergic, beta, receptor kinase 2 /FL=gb:NM_005160.2
203603_s_at	gb:NM_014795.1 /DEF=Homo sapiens zinc finger homeobox 1B (ZFX1B), mRNA. /FEA=mRNA /GEN=ZFX1B /PROD=zinc finger homeobox 1B /DB_XREF=gi:7662183 /UG=Hs:34871 zinc finger homeobox 1B /FL=gb:AB011141.1 gb:NM_014795.1
203786_s_at	gb:NM_003287.1 /DEF=Homo sapiens tumor protein D52-like 1 (TPD52L1), mRNA. /FEA=mRNA /GEN=TPD52L1 /PROD=tumor protein D52-like 1 /DB_XREF=gi:4507640 /UG=Hs:16611 tumor protein D52-like 1 /FL=gb:U44427.1 gb:NM_003287.1

# Affy Probe ID

## Sequence Description

219620\_x\_at gb:NM\_017723.1 /DEF=Homo sapiens hypothetical protein FLJ20245 (FLJ20245), mRNA. /FEA=mRNA /GEN=FLJ20245 /PROD=hypothetical protein FLJ20245 /DB\_XREF=gi:8923220 /UG=Hs.169758 hypothetical protein FLJ20245 /FL=gb:NM\_017723.1

202430\_s\_at gb:NM\_021105.1 /DEF=Homo sapiens phospholipid scramblase 1 (PLSCR1), mRNA. /FEA=mRNA /GEN=PLSCR1 /PROD=phospholipid scramblase 1 /DB\_XREF=gi:10863876 /UG=Hs.198282 phospholipid scramblase 1 /FL=gb:NM\_021105.1 gb:AB006746.1 gb:AF098642.1

217865\_at gb:NM\_018434.1 /DEF=Homo sapiens goliath protein (LOC55819), mRNA. /FEA=mRNA /GEN=LOC55819 /PROD=goliath protein /DB\_XREF=gi:10092650 /UG=Hs.102737 goliath protein /FL=gb:NM\_018434.1

209530\_at gb:U07139.1 /DEF=Human voltage-gated calcium channel beta subunit mRNA, complete cds. /FEA=mRNA /PROD=voltage-gated calcium channel beta subunit /DB\_XREF=gi:463890 /UG=Hs.250712 calcium channel, voltage-dependent, beta 3 subunit /FL=gb:NM\_000725.1 gb:U07139.1

202555\_s\_at gb:NM\_005965.1 /DEF=Homo sapiens myosin, light polypeptide kinase (MYLK), mRNA. /FEA=mRNA /GEN=MYLK /PROD=myosin, light polypeptide kinase /DB\_XREF=gi:5174600 /UG=Hs.211582 myosin, light polypeptide kinase /FL=gb:AB037663.1 gb:NM\_005965.1 gb:AF069601.2

207857\_at gb:NM\_006866.1 /DEF=Homo sapiens leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 (LILRA2), mRNA. /FEA=mRNA /GEN=LILRA2 /PROD=leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /DB\_XREF=gi:5803067 /UG=Hs.94498 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 /FL=gb:U82275.1 gb:NM\_006866.1

212012\_at Consensus includes gb:BF342851 /FEA=EST /DB\_XREF=gi:11289878 /DB\_XREF=est:602015135F1 /CLONE=IMAGE:4150664 /UG=Hs.118893 Melanoma associated gene

209696\_at gb:D26054.1 /DEF=Human mRNA for fructose-1,6-bisphosphatase, complete cds. /FEA=mRNA /PROD=fructose-1,6-bisphosphatase /DB\_XREF=gi:439225 /UG=Hs.574 fructose-1,6-bisphosphatase 1 /FL=gb:M19922.1 gb:L10320.1 gb:D26054.1 gb:D26055.1 gb:D26056.1 gb:NM\_000507.1

202510\_s\_at gb:NM\_006291.1 /DEF=Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA. /FEA=mRNA /GEN=TNFAIP2 /PROD=tumor necrosis factor, alpha-induced protein 2 /DB\_XREF=gi:5454133 /UG=Hs.101382 tumor necrosis factor, alpha-induced protein 2 /FL=gb:M92357.1 gb:NM\_006291.1

214927\_at Consensus includes gb:AL359052.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 1968422. /FEA=mRNA /PROD=ITGBL1, integrin beta-like 1 /DB\_XREF=gi:8518175 /UG=Hs.311054 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 1968422

**Affy Probe ID**

**Sequence Description**

218614\_at gb:NM\_018169.1 /DEF=Homo sapiens hypothetical protein FLJ10652 (FLJ10652), mRNA. /FEA=mRNA /GEN=FLJ10652 /PROD=hypothetical protein FLJ10652 /DB\_XREF=gi:8922572 /UG=Hs.236844 hypothetical protein FLJ10652 /FL=gb:NM\_018169.1

210592\_s\_at gb:M55580.1 /DEF=Human spermidinespermine N1-acetyltransferase mRNA, complete cds. /FEA=mRNA /GEN=spermidinespermine N1-acetyltransferase /PROD=spermidinespermine N1-acetyltransferase /DB\_XREF=gi:338335 /UG=Hs.28491 spermidinespermine N1-acetyltransferase /FL=gb:M55580.1

200636\_s\_at gb:NM\_002840.1 /DEF=Homo sapiens protein tyrosine phosphatase, receptor type, F (PTPRF), mRNA. /FEA=mRNA /GEN=PTPRF /PROD=protein tyrosine phosphatase, receptor type, fpolypeptide /DB\_XREF=gi:4506310 /UG=Hs.75216 protein tyrosine phosphatase, receptor type, F /FL=gb:NM\_002840.1

211075\_s\_at gb:Z25521.1 /DEF=H.sapiens integrin associated protein mRNA, complete CDS,. /FEA=mRNA /PROD=integrin associated protein /DB\_XREF=gi:396704 /FL=gb:Z25521.1

205640\_at gb:NM\_000694.1 /DEF=Homo sapiens aldehyde dehydrogenase 3 family, member B1 (ALDH3B1), mRNA. /FEA=mRNA /GEN=ALDH3B1 /PROD=aldehyde dehydrogenase 3B1 /DB\_XREF=gi:4502042 /UG=Hs.83155 aldehyde dehydrogenase 3 family, member B1 /FL=gb:NM\_000694.1 gb:U10868.1

211138\_s\_at gb:BC005297.1 /DEF=Homo sapiens, Similar to kynurenine 3-monooxygenase (kynurenine 3-hydroxylase), clone MGC:12362, mRNA, complete cds. /FEA=mRNA /PROD=Similar to kynurenine 3-monooxygenase(kynurenine 3-hydroxylase) /DB\_XREF=gi:13529016 /UG=Hs.107318 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) /FL=gb:BC005297.1

204575\_s\_at gb:U38321.1 /DEF=Homo sapiens clone rasi-11 matrix metalloproteinase RASI-1 mRNA, complete cds. /FEA=mRNA /PROD=matrix metalloproteinase RASI-1 /DB\_XREF=gi:2228243 /UG=Hs.154057 matrix metalloproteinase 19 /FL=gb:NM\_002429.2 gb:U38321.1 gb:U37791.1

222237\_s\_at Consensus includes gb:AC084239 /DEF=Homo sapiens chromosome 19, BAC CTC-512J12 (BC347040), complete sequence /FEA=mRNA\_2 /DB\_XREF=gi:10864171 /UG=Hs.48589 zinc finger protein 228

202664\_at Consensus includes gb:AW058622 /FEA=EST /DB\_XREF=gi:5934261 /DB\_XREF=est:wy59c01.x1 /CLONE=IMAGE:2552832 /UG=Hs.24143 Wiskott-Aldrich syndrome protein interacting protein /FL=gb:NM\_003387.2

210241\_s\_at gb:AB007458.1 /DEF=Homo sapiens mRNA for P53TG1-D, complete cds. /FEA=mRNA /GEN=P53TG1 /PROD=P53TG1-D /DB\_XREF=gi:5006272 /UG=Hs.306301 Homo sapiens mRNA for P53TG1-D, complete cds /FL=gb:AB007458.1

Affy Probe_ID	Sequence Description
219183_s_at	gb:NM_013385.2 /DEF=Homo sapiens pleckstrin homology, Sec7 and coiledcoil domains 4 (PSCD4), mRNA. /FEA=mRNA /GEN=PSCD4 /PROD=pleckstrin homology, Sec7 and coiledcoil domains 4 /DB_XREF=gi:8670549 /UG=Hs.7189 pleckstrin homology, Sec7 and coiledcoil domains 4 /FL=gb:AF075458.1 gb:NM_013385.2
206022_at	gb:NM_000266.1 /DEF=Homo sapiens Norrie disease (pseudoglioma) (NDP), mRNA. /FEA=mRNA /GEN=NDP /PROD=Norrie disease protein /DB_XREF=gi:4557788 /UG=Hs.2839 Norrie disease (pseudoglioma) /FL=gb:NM_000266.1
216136_at	Consensus includes gb:AF113683.1 /DEF=Homo sapiens clone FLB3521. /FEA=mRNA /DB_XREF=gi:6855610 /UG=Hs.306569 Homo sapiens clone FLB3521
219059_s_at	Consensus includes gb:AL574194 /FEA=EST /DB_XREF=gi:12934166 /DB_XREF=est:AL574194 /CLONE=CS0DI039YJ06 (3 prime) /UG=Hs.17917 extracellular link domain-containing 1 /FL=gb:AF118108.1 gb:NM_006691.1
211135_x_at	gb:AF009644.1 /DEF=Homo sapiens clone 41 immunoglobulin-like transcript 5 protein mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 5 protein /DB_XREF=gi:2662447 /UG=Hs.105928 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 /FL=gb:AF009644.1
203060_s_at	gb:AF074331.1 /DEF=Homo sapiens PAPS synthetase-2 (PAPSS2) mRNA, complete cds. /FEA=mRNA /GEN=PAPSS2 /PROD=PAPS synthetase-2 /DB_XREF=gi:5052074 /UG=Hs.274230 3-phosphoadenosine 5-phosphosulfate synthase 2 /FL=gb:AF150754.2 gb:AF313907.1 gb:AF091242.1 gb:NM_004670.1 gb:AF074331.1 gb:AF173365.1
34726_at	Cluster Incl. U07139:Human voltage-gated calcium channel beta subunit mRNA, complete cds /cds=(34,1488) /gb=U07139 /gi=463890 /ug=Hs.239447 /len=2532
208370_s_at	gb:NM_004414.2 /DEF=Homo sapiens Down syndrome critical region gene 1 (DSCR1), mRNA. /FEA=mRNA /GEN=DSCR1 /PROD=Down syndrome critical region protein 1 /DB_XREF=gi:7657042 /UG=Hs.184222 Down syndrome critical region gene 1 /FL=gb:U28833.2 gb:NM_004414.2
210132_at	Consensus includes gb:AW189015 /FEA=EST /DB_XREF=gi:6463451 /DB_XREF=est:xk98d10.x1 /CLONE=IMAGE:2674771 /UG=Hs.37054 ephrin-A3 /FL=gb:NM_004952.1 gb:L37360.1 gb:U14187.1
209147_s_at	gb:AB000888.1 /DEF=Homo sapiens mRNA for phosphatidic acid phosphatase 2a, complete cds. /FEA=mRNA /PROD=phosphatidic acid phosphatase 2a /DB_XREF=gi:2467297 /UG=Hs.41569 phosphatidic acid phosphatase type 2A /FL=gb:AB000888.1 gb:AF017116.1 gb:AF014402.1 gb:NM_003711.1

**Affy Probe\_ID****Sequence Description**

212268\_at Consensus includes gb:NM\_030666.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 (SERPINB1), mRNA. /FEA=CDS /GEN=SERPINB1 /PROD=serine (or cysteine) proteinase inhibitor, cladeB (ovalbumin), member 1 /DB\_XREF=gi:13489086 /UG=Hs.183583 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 /FL=gb:NM\_030666.1

220197\_at gb:NM\_020632.1 /DEF=Homo sapiens ATPase, H(+)-transporting, lysosomal, noncatalytic accessory protein 1B (ATP6N1B), mRNA. /FEA=mRNA /GEN=ATP6N1B /PROD=ATPase, H(+)-transporting, lysosomal, noncatalytic accessory protein 1B /DB\_XREF=gi:10190665 /UG=Hs.98967 ATPase, H(+)-transporting, lysosomal, noncatalytic accessory protein 1B /FL=gb:NM\_020632.1 gb:AF245517.1

205614\_x\_at gb:NM\_020998.1 /DEF=Homo sapiens macrophage stimulating 1 (hepatocyte growth factor-like) (MST1), mRNA. /FEA=mRNA /GEN=MST1 /PROD=macrophage stimulating 1 (hepatocyte growthfactor-like) /DB\_XREF=gi:10337614 /UG=Hs.278657 macrophage stimulating 1 (hepatocyte growth factor-like) /FL=gb:NM\_020998.1 gb:M74178.1 gb:L11924.1

213695\_at Consensus includes gb:L48516.1 /DEF=Homo sapiens paraoxonase 3 (PON3) mRNA, 3 end of cds. /FEA=mRNA /GEN=PON3 /PROD=paraoxonase-3 /DB\_XREF=gi:1333633 /UG=Hs.296259 paraoxonase 3

208578\_at gb:NM\_006514.1 /DEF=Homo sapiens sodium channel, voltage-gated, type X, alpha polypeptide (SCN10A), mRNA. /FEA=CDS /GEN=SCN10A /PROD=sodium channel, voltage-gated, type X, alphapolypeptide /DB\_XREF=gi:5730032 /UG=Hs.250443 sodium channel, voltage-gated, type X, alpha polypeptide /FL=gb:AF117907.1 gb:NM\_006514.1

212848\_s\_at Consensus includes gb:BG036668 /FEA=EST /DB\_XREF=gi:12432092 /DB\_XREF=est:602326882F1 /CLONE=IMAGE:4428237 /UG=Hs.18075 chromosome 9 open reading frame 3

210889\_s\_at gb:M31933.1 /DEF=Human IgG low affinity Fc fragment receptor (FcRIIb3) mRNA, complete cds. /FEA=mRNA /DB\_XREF=gi:182598 /UG=Hs.278443 Fc fragment of IgG, low affinity IIb, receptor for (CD32) /FL=gb:M31933.1

210225\_x\_at gb:AF009635.1 /DEF=Homo sapiens clone 17.7 immunoglobulin-like transcript 5 protein mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 5 protein /DB\_XREF=gi:2662429 /UG=Hs.105928 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 /FL=gb:AF000575.1 gb:U91928.1 gb:AF025533.1 gb:AF009632.1 gb:AF009633.1 gb:AF009635.1 gb:AF009638.1 gb:AF009639.1 gb:AF009641.1 gb:AF009642.1 gb:AF031553.1 gb:AF031554.1 gb:AF031555.1 gb:AF031556.1 gb:NM\_006864.1



# Affy Probe\_ID

## Sequence Description

203705_s_at	Consensus includes gb:AI333651 /FEA=EST /DB_XREF=gi:4070210 /DB_XREF=est:qq08e12.x1 /CLONE=IMAGE:1931950 /UG=Hs.173859 frizzled (Drosophila) homolog 7 /FL=gb:AB010881.1 gb:AB017365.1 gb:NM_003507.1
205660_at	gb:NM_003733.1 /DEF=Homo sapiens 2-5oligoadenylate synthetase-like (OASL), mRNA. /FEA=mRNA /GEN=OASL /PROD=2-5oligoadenylate synthetase-like /DB_XREF=gi:11321576 /UG=Hs.118633 2-5oligoadenylate synthetase-like /FL=gb:NM_003733.1 gb:AF063611.1
200906_s_at	Consensus includes gb:AK025843.1 /DEF=Homo sapiens cDNA: FLJ22190 fis, clone HRC01053. /FEA=mRNA /DB_XREF=gi:10438481 /UG=Hs.194431 palladin /FL=gb:AF077041.1 gb:AF151909.1 gb:NM_016081.1
204336_s_at	gb:NM_005873.1 /DEF=Homo sapiens regulator of G-protein signalling 19 (RGS19), mRNA. /FEA=mRNA /GEN=RGS19 /PROD=G protein signalling regulator 19 /DB_XREF=gi:5031704 /UG=Hs.22698 regulator of G-protein signalling 19 /FL=gb:NM_005873.1
216248_s_at	Consensus includes gb:S77154.1 /DEF=TINUR= NGFI-Bnur77 beta-type transcription factor homolog human, T lymphoid cell line, PEER, mRNA, 2469 nt. /FEA=mRNA /GEN=TINUR /DB_XREF=gi:913966 /UG=Hs.82120 nuclear receptor subfamily 4, group A, member 2
217738_at	Consensus includes gb:BF575514 /FEA=EST /DB_XREF=gi:11649318 /DB_XREF=est:602133090F1 /CLONE=IMAGE:4288079 /UG=Hs.239138 pre-B-cell colony-enhancing factor /FL=gb:U02020.1 gb:NM_005746.1
209292_at	Consensus includes gb:AL022726 /DEF=Human DNA sequence from clone 625H18 on chromosome 6p22.2-23. Contains the gene for ID4 Helix-loop-helix DNA binding protein and part of an alternatively spliced novel gene.
216236_s_at	Contains ESTs, STSs, GSSs and putative CpG islands /FEA=mRNA /DB_XREF=gi:3676217 /UG=Hs.34853 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein /FL=gb:NM_001546.1 gb:U16153.1 gb:U28368.1
211990_at	Consensus includes gb:AL110298.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564K1672 (from clone DKFZp564K1672); partial cds. /FEA=mRNA /GEN=DKFZp564K1672 /PROD=hypothetical protein /DB_XREF=gi:5817258 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3
209309_at	Consensus includes gb:M27487.1 /DEF=Homo sapiens MHC class II DPw3-alpha-1 chain mRNA, complete cds. /FEA=CDS /GEN=HLA-DPA1 /PROD=MHC class II DP3-alpha /DB_XREF=gi:703088 /UG=Hs.914 Human mRNA for SB classII histocompatibility antigen alpha-chain /FL=gb:M27487.1
	gb:D90427.1 /DEF=Human mRNA for zinc-alpha2-glycoprotein, complete cds. /FEA=mRNA /PROD=zinc-alpha2-glycoprotein precursor /DB_XREF=gi:220150 /UG=Hs.71 alpha-2-glycoprotein 1, zinc /FL=gb:D90427.1 gb:M76707.1 gb:NM_001185.1

**Affy Probe ID**

**Sequence Description**

208303\_s\_at gb:NM\_022148.1 /DEF=Homo sapiens cytokine receptor-like factor 2 (CRLF2), mRNA. /FEA=mRNA /GEN=CRLF2 /PROD=cytokine receptor-like factor 2 /DB\_XREF=gi:13375623 /UG=Hs.287729 cytokine receptor-like factor 2 /FL=gb:NM\_022148.1

204759\_at gb:NM\_001268.1 /DEF=Homo sapiens chromosome condensation 1-like (CHC1L), mRNA. /FEA=mRNA /GEN=CHC1L /PROD=RCC1-like G exchanging factor RLG /DB\_XREF=gi:4557444 /UG=Hs.27007 chromosome condensation 1-like /FL=gb:AF060219.1 gb:NM\_001268.1

204256\_at gb:NM\_024090.1 /DEF=Homo sapiens hypothetical protein MGC5487 (MGC5487), mRNA. /FEA=mRNA /GEN=MGC5487 /PROD=hypothetical protein MGC5487 /DB\_XREF=gi:13129087 /UG=Hs.211556 hypothetical protein MGC5487 /FL=gb:NM\_024090.1

214974\_x\_at Consensus includes gb:AK026546.1 /DEF=Homo sapiens cDNA: FLJ22893 fis, clone KAT04792. /FEA=mRNA /DB\_XREF=gi:10439427 /UG=Hs.287716 Homo sapiens cDNA: FLJ22893 fis, clone KAT04792

205479\_s\_at gb:NM\_002658.1 /DEF=Homo sapiens plasminogen activator, urokinase (PLAU), mRNA. /FEA=mRNA /GEN=PLAU /PROD=plasminogen activator, urokinase /DB\_XREF=gi:4505862 /UG=Hs.77274 plasminogen activator, urokinase /FL=gb:M15476.1 gb:NM\_002658.1

209815\_at Consensus includes gb:BG054916 /FEA=EST /DB\_XREF=gi:12512119 /DB\_XREF=est:nac92b02.x1 /CLONE=IMAGE:3441723 /UG=Hs.159526 patched (Drosophila) homolog /FL=gb:U43148.1

201818\_at gb:NM\_024830.1 /DEF=Homo sapiens hypothetical protein FLJ12443 (FLJ12443), mRNA. /FEA=mRNA /GEN=FLJ12443 /PROD=hypothetical protein FLJ12443 /DB\_XREF=gi:13376233 /UG=Hs.179882 hypothetical protein FLJ12443 /FL=gb:NM\_024830.1

203678\_at gb:NM\_014967.1 /DEF=Homo sapiens KIAA1018 protein (KIAA1018), mRNA. /FEA=mRNA /GEN=KIAA1018 /PROD=KIAA1018 protein /DB\_XREF=gi:7662449 /UG=Hs.5400 KIAA1018 protein /FL=gb:AB023235.1 gb:NM\_014967.1

218326\_s\_at gb:NM\_018490.1 /DEF=Homo sapiens G protein-coupled receptor 48 (GPR48), mRNA. /FEA=mRNA /GEN=GPR48 /PROD=G protein-coupled receptor 48 /DB\_XREF=gi:8923700 /UG=Hs.160271 G protein-coupled receptor 48 /FL=gb:AF257182.1 gb:NM\_018490.1

211248\_s\_at gb:AF283325.1 /DEF=Homo sapiens chordin variant 3 (CHRD) mRNA, complete cds, alternatively spliced. /FEA=mRNA /GEN=CHRD /PROD=chordin variant 3 /DB\_XREF=gi:11494372 /UG=Hs.166186 chordin /FL=gb:AF283325.1

Affy Probe_ID	Sequence Description
206247_at	gb:NM_005931.1 /DEF=Homo sapiens MHC class I polypeptide-related sequence B (MICB), mRNA. /FEA=mRNA /GEN=MICB /PROD=MHC class I polypeptide-related sequence B /DB_XREF=gi:5174564 /UG=Hs.211580 MHC class I polypeptide-related sequence B /FL=gb:NM_005931.1
214284_s_at	Consensus includes gb:AA022949 /FEA=EST /DB_XREF=gi:1487039 /DB_XREF=est:ze72h02.s1 /CLONE=IMAGE:364563 /UG=Hs.49585 fibroblast growth factor 18
216347_s_at	Consensus includes gb:AK023188.1 /DEF=Homo sapiens cDNA FLJ13126 fis, clone NT2RP3002909, weakly similar to P53-BINDING PROTEIN 2. /FEA=mRNA /DB_XREF=gi:10435002 /UG=Hs.6162 KIAA0771 protein
208309_s_at	gb:NM_006785.1 /DEF=Homo sapiens mucosa associated lymphoid tissue lymphoma translocation gene 1 (MALT1), mRNA. /FEA=mRNA /GEN=MALT1 /PROD=mucosa associated lymphoid tissue lymphomatranslocation gene 1 /DB_XREF=gi:5803077 /UG=Hs.180566 mucosa associated lymphoid tissue lymphoma translocation gene 1 /FL=gb:AF316597.1 gb:AF130356.2 gb:NM_006785.1
207367_at	gb:NM_001676.2 /DEF=Homo sapiens ATPase, H+K+ transporting, nongastric, alpha polypeptide (ATP12A), mRNA. /FEA=mRNA /GEN=ATP12A /PROD=ATPase, H+K+ transporting, nongastric, alhapolypeptide /DB_XREF=gi:10280617 /UG=Hs.1165 ATPase, H+K+ transporting, nongastric, alpha polypeptide /FL=gb:NM_001676.2 gb:U02076.1
201911_s_at	gb:NM_005766.1 /DEF=Homo sapiens FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived) (FARP1), mRNA. /FEA=mRNA /GEN=FARP1 /PROD=FERM, RhoGEF, and pleckstrin domain protein 1 /DB_XREF=gi:5031632 /UG=Hs.183738 FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived) /FL=gb:AB008430.1 gb:NM_005766.1
205101_at	gb:NM_000246.1 /DEF=Homo sapiens MHC class II transactivator (MHC2TA), mRNA. /FEA=mRNA /GEN=MHC2TA /PROD=MHC class II transactivator /DB_XREF=gi:4557748 /UG=Hs.3076 MHC class II transactivator /FL=gb:NM_000246.1 gb:U18259.1
204730_at	gb:NM_014747.1 /DEF=Homo sapiens KIAA0237 gene product (KIAA0237), mRNA. /FEA=mRNA /GEN=KIAA0237 /PROD=KIAA0237 gene product /DB_XREF=gi:7662015 /UG=Hs.78748 KIAA0237 gene product /FL=gb:D87074.1 gb:NM_014747.1
204197_s_at	gb:NM_004350.1 /DEF=Homo sapiens runt-related transcription factor 3 (RUNX3), mRNA. /FEA=mRNA /GEN=RUNX3 /PROD=runt-related transcription factor 3 /DB_XREF=gi:4757917 /UG=Hs.170019 runt-related transcription factor 3 /FL=gb:NM_004350.1

**Affy Probe ID**

**Sequence Description**

204975\_at gb:NM\_001424.1 /DEF=Homo sapiens epithelial membrane protein 2 (EMP2), mRNA. /FEA=mRNA /GEN=EMP2 /PROD=epithelial membrane protein 2 /DB\_XREF=gi:4503560 /UG=Hs.29191 epithelial membrane protein 2 /FL=gb:U52100.1 gb:NM\_001424.1

204622\_x\_at gb:NM\_006186.1 /DEF=Homo sapiens nuclear receptor subfamily 4, group A, member 2 (NR4A2), mRNA. /FEA=mRNA /GEN=NR4A2 /PROD=nuclear receptor subfamily 4, group A, member 2 /DB\_XREF=gi:5453821 /UG=Hs.82120 nuclear receptor subfamily 4, group A, member 2 /FL=gb:NM\_006186.1

212657\_s\_at Consensus includes gb:U65590 /DEF=Homo sapiens IL-1 receptor antagonist IL-1Ra (IL-1RN) gene, alternatively spliced forms, complete cds /FEA=mRNA\_2 /DB\_XREF=gi:2707374 /UG=Hs.81134 interleukin 1 receptor antagonist gb:M17565.1 /DEF=Human MHC class II DQ-beta associated with DRw6, DQw1 protein, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB\_XREF=gi:188188 /FL=gb:M17565.1

211654\_x\_at gb:NM\_021971.1 /DEF=Homo sapiens GDP-mannose pyrophosphorylase B (GMPPB), transcript variant 2, mRNA. /FEA=mRNA /GEN=GMPPB /PROD=GDP-mannose pyrophosphorylase B, isoform 2 /DB\_XREF=gi:11761620 /UG=Hs.28077 GDP-mannose pyrophosphorylase B /FL=gb:NM\_021971.1 gb:BC001141.1 gb:AF135421.1

219920\_s\_at Consensus includes gb:AL023653 /DEF=Human DNA sequence from clone 753P9 on chromosome Xq25-26.1. Contains the gene coding for Aminopeptidase P (EC 3.4.11.9, XAA-ProX-ProProlineAminoacylproline Aminopeptidase) and a novel gene. Contains ESTs, STSs, GSSs and a gaaa repeat polymorphism /FEA=mRNA\_1 /DB\_XREF=gi:3550108 /UG=Hs.61469 hypothetical protein /FL=gb:NM\_018990.1

204923\_at gb:BC005930.1 /DEF=Homo sapiens, Similar to CD58 antigen, (lymphocyte function-associated antigen 3), clone MGC:14538, mRNA, complete cds. /FEA=mRNA /PROD=Similar to CD58 antigen, (lymphocytefunction-associated antigen 3) /DB\_XREF=gi:13543544 /FL=gb:BC005930.1

211744\_s\_at Consensus includes gb:AV717590 /FEA=EST /DB\_XREF=gi:10814742 /DB\_XREF=est:AV717590 /CLONE=DCBCFE01 /UG=Hs.205353 ectonucleoside triphosphate diphosphohydrolase 1 /FL=gb:U87967.1

209473\_at gb:NM\_001406.1 /DEF=Homo sapiens ephrin-B3 (EFNB3), mRNA. /FEA=mRNA /GEN=EFNB3 /PROD=ephrin-B3 precursor /DB\_XREF=gi:4503488 /UG=Hs.26988 ephrin-B3 /FL=gb:U62775.1 gb:U66406.1 gb:NM\_001406.1

205031\_at gb:NM\_017763.1 /DEF=Homo sapiens hypothetical protein FLJ20315 (FLJ20315), mRNA. /FEA=mRNA /GEN=FLJ20315 /PROD=hypothetical protein FLJ20315 /DB\_XREF=gi:8923298 /UG=Hs.18457 hypothetical protein FLJ20315 /FL=gb:NM\_017763.1

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**Sequence Description**

214453\_s\_at Consensus includes gb:NM\_006417.1 /DEF=Homo sapiens interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD) (MTAP44), mRNA. /FEA=CDS /GEN=MTAP44 /PROD=interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD) /DB\_XREF=gi:5453743 /UG=Hs.82316 interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD) /FL=gb:NM\_006417.1

202388\_at gb:NM\_002923.1 /DEF=Homo sapiens regulator of G-protein signalling 2, 24kD (RGS2), mRNA. /FEA=mRNA /GEN=RGS2 /PROD=regulator of G-protein signalling 2, 24kD /DB\_XREF=gi:4506516 /UG=Hs.78944 regulator of G-protein signalling 2, 24kD /FL=gb:L13463.1 gb:NM\_002923.1

218285\_s\_at gb:NM\_020139.1 /DEF=Homo sapiens oxidoreductase UCPA (LOC56898), mRNA. /FEA=mRNA /GEN=LOC56898 /PROD=oxidoreductase UCPA /DB\_XREF=gi:10047131 /UG=Hs.124696 oxidoreductase UCPA /FL=gb:NM\_020139.1 gb:AF164790.1

200905\_x\_at gb:NM\_005516.1 /DEF=Homo sapiens major histocompatibility complex, class I, E (HLA-E), mRNA. /FEA=mRNA /GEN=HLA-E /PROD=major histocompatibility complex, class I, E /DB\_XREF=gi:5031744 /UG=Hs.181392 major histocompatibility complex, class I, E /FL=gb:BC002578.1 gb:NM\_005516.1

200897\_s\_at gb:NM\_016081.1 /DEF=Homo sapiens palladin (KIAA0992), mRNA. /FEA=mRNA /GEN=KIAA0992 /PROD=palladin /DB\_XREF=gi:7706354 /UG=Hs.194431 palladin /FL=gb:AF077041.1 gb:AF151909.1 gb:NM\_016081.1

217456\_x\_at Consensus includes gb:M31183.1 /DEF=Homo sapiens (cosmid clone cd3.3) MHC class I HLA-B51 mRNA, 3 end of cds. /FEA=mRNA /GEN=HLA-B /PROD=MHC class I lymphocyte antigen /DB\_XREF=gi:601851 /UG=Hs.181392 major histocompatibility complex, class I, E

210886\_x\_at gb:AB007457.1 /DEF=Homo sapiens mRNA for P53TG1-C, complete cds. /FEA=mRNA /GEN=P53TG1 /PROD=P53TG1-C /DB\_XREF=gi:5006270 /UG=Hs.274329 TP53 target gene 1 /FL=gb:AB007457.1

221737\_at Consensus includes gb:AK024696.1 /DEF=Homo sapiens cDNA: FLJ21043 fis, clone CAE11633. /FEA=mRNA /DB\_XREF=gi:10437043 /UG=Hs.182874 guanine nucleotide binding protein (G protein) alpha 12 /FL=gb:L01694.1 gb:NM\_007353.1

201012\_at gb:NM\_000700.1 /DEF=Homo sapiens annexin A1 (ANXA1), mRNA. /FEA=mRNA /GEN=ANXA1 /PROD=annexin I /DB\_XREF=gi:4502100 /UG=Hs.78225 annexin A1 /FL=gb:BC001275.1 gb:NM\_000700.1

212062\_at Consensus includes gb:AB014511.1 /DEF=Homo sapiens mRNA for KIAA0611 protein, partial cds. /FEA=mRNA /GEN=KIAA0611 /PROD=KIAA0611 protein /DB\_XREF=gi:3327035 /UG=Hs.70604 ATPase, Class II, type 9A

# **Affy Probe\_ID**

## **Sequence Description**

205833\_s\_at Consensus includes gb:AI770098 /FEA=EST /DB\_XREF=gi:5236553 /DB\_XREF=est:wj30h12.x1 /CLONE=IMAGE:2404391 /UG=Hs.96744 prostate androgen-regulated transcript 1 /FL=gb:NM\_016590.2 gb:AF163475.1

204972\_at gb:NM\_016817.1 /DEF=Homo sapiens 2-5oligoadenylate synthetase 2 (OAS2), transcript variant 1, mRNA. /FEA=mRNA /GEN=OAS2 /PROD=2-5oligoadenylate synthetase 2, isoform p71 /DB\_XREF=gi:8051624 /UG=Hs.264981 2-5oligoadenylate synthetase 2 /FL=gb:M87434.1 gb:NM\_016817.1

217966\_s\_at gb:NM\_022083.1 /DEF=Homo sapiens niban protein (NIBAN), mRNA. /FEA=mRNA /GEN=NIBAN /PROD=niban protein /DB\_XREF=gi:11545796 /UG=Hs.48778 niban protein /FL=gb:AB050477.1 gb:NM\_022083.1 gb:AF288391.1

208158\_s\_at gb:NM\_018030.1 /DEF=Homo sapiens oxysterol-binding protein-related protein 1 (FLJ10217), mRNA. /FEA=mRNA /GEN=FLJ10217 /PROD=oxysterol-binding protein-related protein 1 /DB\_XREF=gi:13877169 /FL=gb:NM\_018030.1

202156\_s\_at Consensus includes gb:N36839 /FEA=EST /DB\_XREF=gi:1157981 /DB\_XREF=est:yy35f07.s1 /CLONE=IMAGE:273253 /UG=Hs.211610 CUG triplet repeat, RNA-binding protein 2 /FL=gb:U69546.1 gb:AF036956.1 gb:AF090694.1 gb:NM\_006561.1

211506\_s\_at gb:AF043337.1 /DEF=Homo sapiens interleukin 8 C-terminal variant (IL8) mRNA, complete cds. /FEA=mRNA /GEN=IL8 /PROD=interleukin 8 C-terminal variant /DB\_XREF=gi:12641914 /UG=Hs.624 interleukin 8 /FL=gb:AF043337.1

209153\_s\_at gb:M31523.1 /DEF=Human transcription factor (E2A) mRNA, complete cds. /FEA=mRNA /GEN=TCF3 /DB\_XREF=gi:339477 /UG=Hs.101047 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12E47) /FL=gb:M31523.1

212730\_at Consensus includes gb:AK026420.1 /DEF=Homo sapiens cDNA: FLJ22767 fis, clone KAI1191. /FEA=mRNA /DB\_XREF=gi:10439281 /UG=Hs.10587 KIAA0353 protein

211404\_s\_at gb:BC004371.1 /DEF=Homo sapiens, clone MGC:10449, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10449) /DB\_XREF=gi:13325115 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC004371.1

215017\_s\_at Consensus includes gb:AW270932 /FEA=EST /DB\_XREF=gi:6657962 /DB\_XREF=est:xs06c12.x1 /CLONE=IMAGE:2768854 /UG=Hs.239681 hypothetical protein FLJ20275

1007\_s\_at U48705 /FEATURE=mRNA /DEFINITION=HSU48705 Human receptor tyrosine kinase DDR gene, complete cds

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221900_at	Consensus includes gb:AI806793 /FEA=EST /DB_XREF=gi:5393359 /DB_XREF=est:wfl5d05.x1 /CLONE=IMAGE:2350665 /UG=Hs.249239 collagen, type VIII, alpha 2
205872_x_at	gb:NM_022359.1 /DEF=Homo sapiens similar to rat myomegalin (LOC64182), mRNA. /FEA=mRNA /GEN=LOC64182 /PROD=similar to rat myomegalin /DB_XREF=gi:11641276 /UG=Hs.265848 similar to rat myomegalin /FL=gb:AB042558.1 gb:NM_022359.1
210517_s_at	gb:AB003476.1 /DEF=Homo sapiens mRNA for gravin, complete cds. /FEA=mRNA /PROD=gravin /DB_XREF=gi:2081606 /UG=Hs.788 A kinase (PRKA) anchor protein (gravin) 12 /FL=gb:AB003476.1
214574_x_at	Consensus includes gb:NM_007161.1 /DEF=Homo sapiens DNA segment on chromosome 6 (unique) 49 expressed sequence, NK cell triggering receptor, p30 (D6S49E), mRNA. /FEA=CDS /GEN=D6S49E /PROD=leukocyte-specific transcript 1 /DB_XREF=gi:6005740 /UG=Hs.88411 lymphocyte antigen 117 /FL=gb:NM_007161.1
202944_at	gb:NM_000262.1 /DEF=Homo sapiens N-acetylgalactosaminidase, alpha- (NAGA), mRNA. /FEA=mRNA /GEN=NAGA /PROD=alpha-N-acetylgalactosaminidase precursor /DB_XREF=gi:4557780 /UG=Hs.75372 N-acetylgalactosaminidase, alpha- /FL=gb:BC000095.1 gb:M62783.1 gb:M38083.1 gb:NM_000262.1
210146_x_at	gb:AF004231.1 /DEF=Homo sapiens monocytemacrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds. /FEA=mRNA /GEN=MIR cl-10 /PROD=MIR-10 /DB_XREF=gi:2343110 /UG=Hs.22405 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2 /FL=gb:AF000574.1 gb:AF004231.1 gb:AF011565.1
217997_at	Consensus includes gb:AI795908 /FEA=EST /DB_XREF=gi:5361371 /DB_XREF=est:wh40a05.x1 /CLONE=IMAGE:2383184 /UG=Hs.82101 pleckstrin homology-like domain, family A, member 1 /FL=gb:NM_007350.1
220330_s_at	gb:NM_022136.1 /DEF=Homo sapiens SAM domain, SH3 domain and nuclear localisation signals, 1 (SAMSNI), mRNA. /FEA=mRNA /GEN=SAMSNI /PROD=SAM domain, SH3 domain and nuclear localisationsignals, 1 /DB_XREF=gi:11545870 /UG=Hs.24633 SAM domain, SH3 domain and nuclear localisation signals, 1 /FL=gb:AF222927.1 gb:NM_022136.1
206114_at	gb:NM_004438.1 /DEF=Homo sapiens EphA4 (EPhA4), mRNA. /FEA=mRNA /GEN=EPhA4 /PROD=EPhA4 /DB_XREF=gi:4758279 /UG=Hs.73964 EphA4 /FL=gb:NM_004438.1 gb:L36645.1
200907_s_at	Consensus includes gb:AU157932 /FEA=EST /DB_XREF=gi:11019453 /DB_XREF=est:AU157932 /CLONE=PLACE1010217 /UG=Hs.194431 palladin /FL=gb:AF077041.1 gb:AF151909.1 gb:NM_016081.1

Affy Probe_ID	Sequence Description
214068_at	Consensus includes gb:AF070610.1 /DEF=Homo sapiens clone 24505 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387992 /UG=Hs.100543 Homo sapiens clone 24505 mRNA sequence
212372_at	Consensus includes gb:AK026977.1 /DEF=Homo sapiens cDNA: FLJ23324 fis, clone HEP12482, highly similar to HUMMYOHC Human nonmuscle myosin heavy chain-B (MYH10) mRNA. /FEA=mRNA /DB_XREF=gi:10439970 /UG=Hs.296842 Homo sapiens, clone IMAGE:3357927, mRNA, partial cds
202504_at	gb:NM_012101.1 /DEF=Homo sapiens ataxia-telangiectasia group D-associated protein (ATDC), mRNA. /FEA=mRNA /GEN=ATDC /PROD=ataxia-telangiectasia group D-associated protein /DB_XREF=gi:6912249 /UG=Hs.82237 ataxia-telangiectasia group D-associated protein /FL=gb:AF230388.1 gb:L24203.1 gb:NM_012101.1
213187_x_at	Consensus includes gb:BG538564 /FEA=EST /DB_XREF=gi:13530797 /DB_XREF=est:602567289F1 /CLONE=IMAGE:4691639 /UG=Hs.324746 alpha-2-HS-glycoprotein
206873_at	gb:NM_001215.1 /DEF=Homo sapiens carbonic anhydrase VI (CA6), mRNA. /FEA=mRNA /GEN=CA6 /PROD=carbonic anhydrase VI precursor /DB_XREF=gi:4557396 /UG=Hs.100322 carbonic anhydrase VI /FL=gb:M57892.1 gb:NM_001215.1
200798_x_at	gb:NM_021960.1 /DEF=Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1), mRNA. /FEA=mRNA /GEN=MCL1 /PROD=myeloid cell leukemia sequence 1 (BCL2-related) /DB_XREF=gi:11386164 /UG=Hs.86386 myeloid cell leukemia sequence 1 (BCL2-related) /FL=gb:NM_021960.1 gb:AF118124.1
213418_at	Consensus includes gb:NM_002155.1 /DEF=Homo sapiens heat shock 70kD protein 6 (HSP70B) (HSPA6), mRNA. /FEA=CDS /GEN=HSPA6 /PROD=heat shock 70kD protein 6 (HSP70B) /DB_XREF=gi:4504514 /UG=Hs.3268 heat shock 70kD protein 6 (HSP70B) /FL=gb:NM_002155.1
205896_at	gb:NM_003059.1 /DEF=Homo sapiens solute carrier family 22 (organic cation transporter), member 4 (SLC22A4), mRNA. /FEA=mRNA /GEN=SLC22A4 /PROD=solute carrier family 22 (organic cation transporter), member 4 /DB_XREF=gi:4507002 /UG=Hs.77239 solute carrier family 22 (organic cation transporter), member 4 /FL=gb:AB007448.1 gb:NM_003059.1
214607_at	Consensus includes gb:AW085556 /FEA=EST /DB_XREF=gi:6040708 /DB_XREF=est:wy67b11.x1 /CLONE=IMAGE:2553597 /UG=Hs.152663 p21 (CDKN1A)-activated kinase 3 /FL=gb:AF068864.1 gb:NM_002578.1
213100_at	Consensus includes gb:AA127885 /FEA=EST /DB_XREF=gi:1687192 /DB_XREF=est:z113b04.s1 /CLONE=IMAGE:501775 /UG=Hs.13350 Homo sapiens mRNA; cDNA DKFZp586D0918 (from clone DKFZp586D0918)



**Affy Probe ID****Sequence Description**

212509\_s\_at Consensus includes gb:BF968134 /FEA=EST /DB\_XREF=gi:12335349 /DB\_XREF=est:602269121F1 /CLONE=IMAGE:4357349 /UG=Hs.250723 FK506 binding protein 12-rapamycin associated protein 1 gb:Nm\_002167.1 /DEF=Homo sapiens inhibitor of DNA binding 3, dominant negative helix-loop-helix protein (ID3), mRNA. /FEA=mRNA /GEN=ID3 /PROD=inhibitor of DNA binding 3, dominant negative helix-loop-helix protein /DB\_XREF=gi:10835060 /UG=Hs.76884 inhibitor of DNA binding 3, dominant negative helix-loop-helix protein /FL=gb:Nm\_002167.1

209757\_s\_at gb:BC002712.1 /DEF=Homo sapiens, v-myc avian myelocytomatosis viral related oncogene, neuroblastoma derived, clone MGC:3962, mRNA, complete cds. /FEA=mRNA /PROD=v-myc avian myelocytomatosis viral related oncogene, neuroblastoma derived /DB\_XREF=gi:12803748 /UG=Hs.25960 v-myc avian myelocytomatosis viral related oncogene, neuroblastoma derived /FL=gb:BC002712.1 gb:Nm\_005378.1

205151\_s\_at gb:Nm\_014817.1 /DEF=Homo sapiens KIAA0644 gene product (KIAA0644), mRNA. /FEA=mRNA /GEN=KIAA0644 /PROD=KIAA0644 gene product /DB\_XREF=gi:7662219 /UG=Hs.21572 KIAA0644 gene product /FL=gb:AB014544.1 gb:Nm\_014817.1

212286\_at Consensus includes gb:AW572909 /FEA=EST /DB\_XREF=gi:7237642 /DB\_XREF=est:hfl7d03.x1 /CLONE=IMAGE:2932133 /UG=Hs.27973 KIAA0874 protein /FL=gb:AF317425.1

215076\_s\_at Consensus includes gb:AU144167 /FEA=EST /DB\_XREF=gi:11005688 /DB\_XREF=est:AU144167 /CLONE=HEMBA1001071 /UG=Hs.297909 Homo sapiens cDNA FLJ11428 fis, clone HEMBA1001071, highly similar to PROCOLLAGEN ALPHA 1(III) CHAIN PRECURSOR

201298\_s\_at gb:BC003398.1 /DEF=Homo sapiens, hypothetical protein FLJ10788, clone MGC:4929, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ10788 /DB\_XREF=gi:13097287 /UG=Hs.196437 hypothetical protein FLJ10788 /FL=gb:AB016839.1 gb:BC003398.1 gb:Nm\_018221.1

216988\_s\_at Consensus includes gb:L48722 /DEF=Homo sapiens (clone hhl8) protein tyrosine phosphatase (ptp-IV1r) gene, 5 end of cds /FEA=CDS /DB\_XREF=gi:1246237 /UG=Hs.82911 protein tyrosine phosphatase type IVA, member 2

221903\_s\_at Consensus includes gb:BE046443 /FEA=EST /DB\_XREF=gi:8363496 /DB\_XREF=est:hn47d10.x2 /CLONE=IMAGE:3026803 /UG=Hs.18827 KIAA0849 protein

213797\_at Consensus includes gb:AI337069 /FEA=EST /DB\_XREF=gi:4073996 /DB\_XREF=est:qx82h04.x1 /CLONE=IMAGE:2009047 /UG=Hs.17518 Homo sapiens cig5 mRNA, partial sequence

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**Sequence Description**

204393\_s\_at gb:NM\_001099.2 /DEF=Homo sapiens acid phosphatase, prostate (ACPP), mRNA. /FEA=mRNA /GEN=ACPP /PROD=prostatic acid phosphatase precursor /DB\_XREF=gi:6382063 /UG=Hs.1852 acid phosphatase, prostate /FL=gb:M24902.1 gb:M34840.1 gb:NM\_001099.2

212681\_at Consensus includes gb:AI770004 /FEA=EST /DB\_XREF=gi:5236513 /DB\_XREF=est:wh66d06.x1 /CLONE=IMAGE:2385707 /UG=Hs.103839 erythrocyte membrane protein band 4.1-like 3

205119\_s\_at gb:NM\_002029.1 /DEF=Homo sapiens formyl peptide receptor 1 (FPR1), mRNA. /FEA=mRNA /GEN=FPR1 /PROD=formyl peptide receptor 1 /DB\_XREF=gi:4503778 /UG=Hs.753 formyl peptide receptor 1 /FL=gb:BC005315.1 gb:M60626.1 gb:M60627.1 gb:NM\_002029.1

205329\_s\_at gb:NM\_003794.1 /DEF=Homo sapiens sorting nexin 4 (SNX4), mRNA. /FEA=mRNA /GEN=SNX4 /PROD=sorting nexin 4 /DB\_XREF=gi:4507144 /UG=Hs.267812 sorting nexin 4 /FL=gb:AF130078.1 gb:AF065485.1 gb:NM\_003794.1

212013\_at Consensus includes gb:D86983.1 /DEF=Human mRNA for KIAA0230 gene, partial cds. /FEA=mRNA /GEN=KIAA0230 /DB\_XREF=gi:1504039 /UG=Hs.118893 Melanoma associated gene

207114\_at gb:NM\_025261.1 /DEF=Homo sapiens G6C protein (G6C), mRNA. /FEA=mRNA /GEN=G6C /PROD=G6C protein /DB\_XREF=gi:13376873 /UG=Hs.241586 G6C protein /FL=gb:NM\_025261.1

204806\_x\_at gb:NM\_018950.1 /DEF=Homo sapiens major histocompatibility complex, class I, F (HLA-F), mRNA. /FEA=mRNA /GEN=HLA-F /PROD=major histocompatibility complex, class I, F /DB\_XREF=gi:9665231 /UG=Hs.110309 major histocompatibility complex, class I, F /FL=gb:NM\_018950.1

210880\_s\_at gb:AB001467.1 /DEF=Homo sapiens mRNA for Efs2, complete cds. /FEA=mRNA /PROD=Efs2 /DB\_XREF=gi:2829303 /UG=Hs.24587 signal transduction protein (SH3 containing) /FL=gb:AB001467.1

221127\_s\_at gb:NM\_006394.1 /DEF=Homo sapiens regulated in glioma (RIG), mRNA. /FEA=mRNA /GEN=RIG /PROD=regulated in glioma /DB\_XREF=gi:5454007 /UG=Hs.278503 regulated in glioma /FL=gb:NM\_006394.1 gb:U32331.1

212913\_at Consensus includes gb:BE674960 /FEA=EST /DB\_XREF=gi:10035501 /DB\_XREF=est:7f11a09.x1 /CLONE=IMAGE:3294328 /UG=Hs.112193 mutS (E. coli) homolog 5

209890\_at gb:AF065389.1 /DEF=Homo sapiens tetraspan NET-4 mRNA, complete cds. /FEA=mRNA /PROD=tetraspan NET-4 /DB\_XREF=gi:3152702 /UG=Hs.20709 tetraspan 5 /FL=gb:AF065389.1 gb:NM\_005723.1

212820\_at Consensus includes gb:AB020663.1 /DEF=Homo sapiens mRNA for KIAA0856 protein, partial cds. /FEA=mRNA /GEN=KIAA0856 /PROD=KIAA0856 protein /DB\_XREF=gi:4240200 /UG=Hs.13264 KIAA0856 protein

**Affy Probe ID**

**Sequence Description**

210749_x_at	gb:L11315.1 /DEF=Homo sapiens receptor tyrosine kinase mRNA, complete cds. /FEA=mRNA /PROD=receptor tyrosine kinase /DB_XREF=gi:403386 /UG=Hs.75562 discoidin domain receptor family, member 1 /FL=gb:L11315.1
211535_s_at	gb:M60485.1 /DEF=Human fibroblast growth factor receptor mRNA, complete cds. /FEA=mRNA /PROD=fibroblast growth factor receptor /DB_XREF=gi:182560 /UG=Hs.748 fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome) /FL=gb:M60485.1
203373_at	gb:NM_003877.1 /DEF=Homo sapiens STAT induced STAT inhibitor-2 (STATI2), mRNA. /FEA=mRNA /GEN=STATI2 /PROD=STAT induced STAT inhibitor-2 /DB_XREF=gi:4507262 /UG=Hs.110776 STAT induced STAT inhibitor-2 /FL=gb:AB004903.1 gb:AB006966.1 gb:AF037989.1 gb:AF020590.1 gb:NM_003877.1
203413_at	gb:NM_006159.1 /DEF=Homo sapiens nel (chicken)-like 2 (NELL2), mRNA. /FEA=mRNA /GEN=NELL2 /PROD=nel (chicken)-like 2 /DB_XREF=gi:5453765 /UG=Hs.79389 nel (chicken)-like 2 /FL=gb:D83018.1 gb:NM_006159.1
204547_at	gb:NM_006822.1 /DEF=Homo sapiens GTP-binding protein homologous to Saccharomyces cerevisiae SEC4 (SEC4L), mRNA. /FEA=mRNA /GEN=SEC4L /PROD=GTP-binding protein homologous to Saccharomyces cerevisiae SEC4 /DB_XREF=gi:5803162 /UG=Hs.302498 GTP-binding protein homologous to Saccharomyces cerevisiae SEC4 /FL=gb:U05227.1 gb:NM_006822.1
217755_at	gb:NM_016185.1 /DEF=Homo sapiens hematological and neurological expressed 1 (HN1), mRNA. /FEA=mRNA /GEN=HN1 /PROD=hematological and neurological expressed 1 /DB_XREF=gi:7705876 /UG=Hs.109706 hematological and neurological expressed 1 /FL=gb:AF060925.1 gb:BC001420.1 gb:AF177862.1 gb:NM_016185.1
216320_x_at	Consensus includes gb:U37055 /DEF=Human hepatocyte growth factor-like protein gene, complete cds /FEA=mRNA /DB_XREF=gi:1311660 /UG=Hs.278657 macrophage stimulating 1 (hepatocyte growth factor-like)
204134_at	gb:NM_002599.1 /DEF=Homo sapiens phosphodiesterase 2A, cGMP-stimulated (PDE2A), mRNA. /FEA=mRNA /GEN=PDE2A /PROD=phosphodiesterase 2A, cGMP-stimulated /DB_XREF=gi:4505656 /UG=Hs.154437 phosphodiesterase 2A, cGMP-stimulated /FL=gb:U67733.1 gb:NM_002599.1
212593_s_at	Consensus includes gb:N92498 /FEA=EST /DB_XREF=gi:1264807 /DB_XREF=est:zb28a04.s1 /CLONE=IMAGE:304878 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, clone HEP11691
203455_s_at	gb:NM_002970.1 /DEF=Homo sapiens spermidinespermine N1-acetyltransferase (SAT), mRNA. /FEA=mRNA /GEN=SAT /PROD=spermidinespermine N1-acetyltransferase /DB_XREF=gi:4506788 /UG=Hs.28491 spermidinespermine N1-acetyltransferase /FL=gb:BC002503.1 gb:M77693.1 gb:NM_002970.1

Affy Probe_ID	Sequence Description
203765_at	gb:NM_012198.1 /DEF=Homo sapiens grancalcin (GCL), mRNA. /FEA=mRNA /GEN=GCL /PROD=grancalcin /DB_XREF=gi:6912387 /UG=Hs.79381 grancalcin /FL=gb:BC005214.1 gb:M81637.1 gb:NM_012198.1
212594_at	Consensus includes gb:AI185160 /FEA=EST /DB_XREF=gi:3735798 /DB_XREF=est:qe30g10.sl /CLONE=IMAGE:1740546 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, clone HEP11691
205933_at	gb:NM_015559.1 /DEF=Homo sapiens KIAA0437 protein (KIAA0437), mRNA. /FEA=mRNA /GEN=KIAA0437 /PROD=KIAA0437 protein /DB_XREF=gi:7662121 /UG=Hs.151717 KIAA0437 protein /FL=gb:AB022660.1
207169_x_at	gb:NM_015559.1
207169_x_at	gb:NM_001954.2 /DEF=Homo sapiens discoidin domain receptor family, member 1 (DDR1), transcript variant 2, mRNA. /FEA=mRNA /GEN=DDR1 /PROD=discoidin receptor tyrosine kinase isoform b /DB_XREF=gi:7669486 /UG=Hs.75562 discoidin domain receptor family, member 1 /FL=gb:NM_001954.2
212761_at	Consensus includes gb:AI949687 /FEA=EST /DB_XREF=gi:5741997 /DB_XREF=est:wq13g04.x1 /CLONE=IMAGE:2471190 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG-box)
222368_at	Consensus includes gb:AW972351 /FEA=EST /DB_XREF=gi:8162197 /DB_XREF=est:EST384442 /UG=Hs.293451 ESTs
202551_s_at	Consensus includes gb:BG546884 /FEA=EST /DB_XREF=gi:13545549 /DB_XREF=est:602574066F1 /CLONE=IMAGE:4702049 /UG=Hs.19280 cysteine-rich motor neuron 1 /FL=gb:NM_016441.1 gb:AF167706.1
206749_at	gb:NM_001764.1 /DEF=Homo sapiens CD1B antigen, b polypeptide (CD1B), mRNA. /FEA=mRNA /GEN=CD1B /PROD=CD1B antigen, b polypeptide /DB_XREF=gi:4502644 /UG=Hs.1310 CD1B antigen, b polypeptide /FL=gb:M28826.1 gb:NM_001764.1
207808_s_at	gb:NM_000313.1 /DEF=Homo sapiens protein S (alpha) (PROS1), mRNA. /FEA=mRNA /GEN=PROS1 /PROD=protein S (alpha) /DB_XREF=gi:4506116 /UG=Hs.64016 protein S (alpha) /FL=gb:M15036.1
203964_at	gb:NM_000313.1
203964_at	gb:NM_004688.1 /DEF=Homo sapiens N-myc (and STAT) interactor (NMI), mRNA. /FEA=mRNA /GEN=NMI /PROD=N-myc and STAT interactor /DB_XREF=gi:4758813 /UG=Hs.54483 N-myc (and STAT) interactor /FL=gb:BC001268.1 gb:U32849.1 gb:NM_004688.1
207528_s_at	gb:NM_014331.1 /DEF=Homo sapiens solute carrier family 7, (cationic amino acid transporter, y+ system) member 11 (SLC7A11), mRNA. /FEA=mRNA /GEN=SLC7A11 /PROD=solute carrier family 7, (cationic amino acid transporter, y+ system) member 11 /DB_XREF=gi:7657682 /UG=Hs.6682 solute carrier family 7, (cationic amino acid transporter, y+ system) member 11 /FL=gb:AF200708.1 gb:AB026891.1 gb:NM_014331.1

# Affy Probe ID

## Sequence Description

203408\_s\_at gb:NM\_002971.1 /DEF=Homo sapiens special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) (SATB1), mRNA. /FEA=mRNA /GEN=SATB1 /PROD=special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) /DB\_XREF=gi:4506790 /UG=Hs.74592 special AT-rich sequence binding protein 1 (binds to nuclear matrixscaffold-associating DNAs) /FL=gb:M97287.1 gb:NM\_002971.1

207387\_s\_at gb:NM\_000167.1 /DEF=Homo sapiens glycerol kinase (GK), mRNA. /FEA=mRNA /GEN=GK /PROD=glycerol kinase /DB\_XREF=gi:4504006 /UG=Hs.1466 glycerol kinase /FL=gb:L13943.1 gb:NM\_000167.1

207843\_x\_at gb:NM\_001914.1 /DEF=Homo sapiens cytochrome b-5 (CYB5), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=CYB5 /PROD=cytochrome b-5 /DB\_XREF=gi:4503182 /UG=Hs.83834 cytochrome b-5 /FL=gb:M60174.1 gb:NM\_001914.1

217996\_at Consensus includes gb:AA576961 /FEA=EST /DB\_XREF=gi:2354435 /DB\_XREF=est:nm82d08.s1 /CLONE=IMAGE:1074735 /UG=Hs.82101 pleckstrin homology-like domain, family A, member 1 /FL=gb:NM\_007350.1

203657\_s\_at gb:NM\_003793.2 /DEF=Homo sapiens cathepsin F (CTSF), mRNA. /FEA=mRNA /GEN=CTSF /PROD=cathepsin F /DB\_XREF=gi:6042195 /UG=Hs.11590 cathepsin F /FL=gb:AF071748.1 gb:AF071749.1 gb:AF088886.2

208763\_s\_at gb:NM\_003793.2 gb:AF136279.1

220403\_s\_at gb:AL110191.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566A093 (from clone DKFZp566A093); complete cds. /FEA=mRNA /GEN=DKFZp566A093 /PROD=hypothetical protein /DB\_XREF=gi:5817105 /UG=Hs.75450 delta sleep inducing peptide, immunoreactor /FL=gb:AF228339.1 gb:AF153603.1 gb:AL110191.1 gb:AF183393.1

220403\_s\_at gb:NM\_022112.1 /DEF=Homo sapiens p53-regulated apoptosis-inducing protein 1 (P53AIP1), mRNA. /FEA=mRNA /GEN=P53AIP1 /PROD=p53-regulated apoptosis-inducing protein 1 /DB\_XREF=gi:11545826 /UG=Hs.160953 p53-regulated apoptosis-inducing protein 1 /FL=gb:AB045832.1 gb:NM\_022112.1

201876\_at gb:NM\_000305.1 /DEF=Homo sapiens paraoxonase 2 (PON2), mRNA. /FEA=mRNA /GEN=PON2 /PROD=paraoxonase 2 /DB\_XREF=gi:4505952 /UG=Hs.169857 paraoxonase 2 /FL=gb:L48513.1 gb:AF001601.1

221011\_s\_at gb:NM\_000305.1

208702\_x\_at gb:NM\_030915.1 /DEF=Homo sapiens hypothetical protein DKFZp566J091 (DKFZp566J091), mRNA. /FEA=mRNA /GEN=DKFZp566J091 /PROD=hypothetical protein DKFZp566J091 /DB\_XREF=gi:13569871 /FL=gb:NM\_030915.1

208702\_x\_at Consensus includes gb:AI525212 /FEA=EST /DB\_XREF=gi:4439347 /DB\_XREF=est:pt1.1-2.A08.r /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2 /FL=gb:BC000373.1

# Affy Probe\_ID

## Sequence Description

61734_at	Cluster Incl. AI797684:we90d02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2348355 /clone_end=3 /gb=AI797684 /gi=5363156 /ug=Hs.39619 /len=708
209656_s_at	gb:AL136550.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761J17121 (from clone DKFZp761J17121); complete cds. /FEA=mRNA /GEN=DKFZp761J17121 /PROD=hypothetical protein /DB_XREF=gi:13276606 /UG=Hs.8769 Homo sapiens mRNA; cDNA DKFZp761J17121 (from clone DKFZp761J17121); complete cds /FL=gb:AL136550.1 gb:Nm_020386.1 /DEF=Homo sapiens H-REV107 protein-related protein (LOC57110), mRNA. /FEA=mRNA /GEN=LOC57110 /PROD=H-REV107 protein-related protein /DB_XREF=gi:9966858 /UG=Hs.36761 H-REV107 protein-related protein /FL=gb:AB030816.1 gb:Nm_020386.1
219983_at	gb:Nm_002056.1 /DEF=Homo sapiens glutamine-fructose-6-phosphate transaminase 1 (GFPT1), mRNA. /FEA=mRNA /GEN=GFPT1 /PROD=glutamine-fructose-6-phosphate aminotransferase /DB_XREF=gi:4503980 /UG=Hs.1674 glutamine-fructose-6-phosphate transaminase 1 /FL=gb:M90516.1 gb:Nm_002056.1
202722_s_at	gb:AF241787.1 /DEF=Homo sapiens CGI16-iso mRNA, complete cds. /FEA=mRNA /PROD=CGI16-iso /DB_XREF=gi:12005490 /UG=Hs.18625 Mitochondrial Acyl-CoA Thioesterase /FL=gb:AF241787.1 gb:U86214.1 /DEF=Human Fas-associated death domain protein interleukin-1b-converting enzyme 2 mRNA, complete cds. /FEA=mRNA /PROD=Fas-associated death domain protein interleukin-1b-converting enzyme 2 /DB_XREF=gi:1835778 /UG=Hs.5353 caspase 10, apoptosis-related cysteine protease /FL=gb:U86214.1 Consensus includes gb:BE856376 /FEA=EST /DB_XREF=gi:10369339 /DB_XREF=est:7f93f09.x1 /CLONE=IMAGE:3304553 /UG=Hs.277543 very long-chain acyl-CoA synthetase; lipidosin /FL=gb:Nm_015162.1 gb:AF179481.1
210955_at	gb:Nm_004393.1 /DEF=Homo sapiens dystroglycan 1 (dystrophin-associated glycoprotein 1) (DAG1), mRNA. /FEA=mRNA /GEN=DAG1 /PROD=dystroglycan 1 precursor /DB_XREF=gi:4758115 /UG=Hs.76111 dystroglycan 1 (dystrophin-associated glycoprotein 1) /FL=gb:L19711.1 gb:Nm_004393.1
205417_s_at	Consensus includes gb:AW575374 /FEA=EST /DB_XREF=gi:7246913 /DB_XREF=est:UL-HF-BL0-adc-c-12-0-UI.s1 /CLONE=IMAGE:3060887 /UG=Hs.288555 Homo sapiens cDNA: FLJ22425 fis, clone HRC08686
221773_at	Consensus includes gb:AL353949.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761P1114 (from clone DKFZp761P1114). /FEA=mRNA /DB_XREF=gi:7669990 /UG=Hs.306506 Homo sapiens mRNA; cDNA DKFZp761P1114 (from clone DKFZp761P1114)
216704_at	Consensus includes gb:BF511231 /FEA=EST /DB_XREF=gi:11594529 /DB_XREF=est:UL-H-BI4-aoi-g-11-0-UI.s1 /CLONE=IMAGE:3085244 /UG=Hs.288582 ESTs, Weakly similar to ubiquitous TPR motif, Y isoform H.sapiens
213258_at	

# Affy Probe\_ID

## Sequence Description

201525\_at gb:NM\_001647.1 /DEF=Homo sapiens apolipoprotein D (APOD), mRNA. /FEA=mRNA /GEN=APOD /PROD=apolipoprotein D precursor /DB\_XREF=gi:4502162 /UG=Hs.75736 apolipoprotein D /FL=gb:J02611.1 gb:NM\_001647.1

201137\_s\_at gb:NM\_002121.1 /DEF=Homo sapiens major histocompatibility complex, class II, DP beta 1 (HLA-DPB1), mRNA. /FEA=mRNA /GEN=HLA-DPB1 /PROD=major histocompatibility complex, class II, DPbeta 1 /DB\_XREF=gi:4504404 /UG=Hs.814 major histocompatibility complex, class II, DP beta 1 /FL=gb:J03041.1 gb:M57466.1 gb:M83664.1 gb:NM\_002121.1 gb:M28200.1 gb:M28202.1

205968\_at gb:NM\_002252.1 /DEF=Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 (KCNS3), mRNA. /FEA=mRNA /GEN=KCNS3 /PROD=potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 /DB\_XREF=gi:4504862 /UG=Hs.47584 potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 /FL=gb:BC004148.1 gb:BC004987.1 gb:AF043472.1 gb:NM\_002252.1

205569\_at gb:NM\_014398.1 /DEF=Homo sapiens similar to lysosome-associated membrane glycoprotein (TSC403), mRNA. /FEA=mRNA /GEN=TSC403 /PROD=similar to lysosome-associated membraneglycoprotein /DB\_XREF=gi:7657660 /UG=Hs.10887 similar to lysosome-associated membrane glycoprotein /FL=gb:AB013924.1 gb:NM\_014398.1

203355\_s\_at gb:NM\_015310.1 /DEF=Homo sapiens KIAA0942 protein (KIAA0942), mRNA. /FEA=mRNA /GEN=KIAA0942 /PROD=KIAA0942 protein /DB\_XREF=gi:7662395 /UG=Hs.6763 KIAA0942 protein /FL=gb:AF243495.2 gb:NM\_015310.1

206595\_at gb:NM\_001323.1 /DEF=Homo sapiens cystatin EM (CST6), mRNA. /FEA=mRNA /GEN=CST6 /PROD=cystatin M /DB\_XREF=gi:4503112 /UG=Hs.83393 cystatin EM /FL=gb:U62800.1 gb:U81233.1 gb:NM\_001323.1

212186\_at Consensus includes gb:BE855983 /FEA=EST /DB\_XREF=gi:10368561 /DB\_XREF=est:7f85g11.x1 /CLONE=IMAGE:3303812 /UG=Hs.7232 acetyl-Coenzyme A carboxylase alpha /FL=gb:NM\_000664.1 gb:U19822.1

214259\_s\_at Consensus includes gb:AI144075 /FEA=EST /DB\_XREF=gi:3665884 /DB\_XREF=est:qi63e12.x1 /CLONE=IMAGE:1861198 /UG=Hs.6980 aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)

204787\_at gb:NM\_007268.1 /DEF=Homo sapiens Ig superfamily protein (Z39IG), mRNA. /FEA=mRNA /GEN=Z39IG /PROD=Ig superfamily protein /DB\_XREF=gi:6005957 /UG=Hs.8904 Ig superfamily protein /FL=gb:NM\_007268.1

209014\_at gb:AF217963.1 /DEF=Homo sapiens NRAGE mRNA, complete cds. /FEA=mRNA /PROD=NRAGE /DB\_XREF=gi:9963809 /UG=Hs.177556 melanoma antigen, family D, 1 /FL=gb:AF132205.1 gb:AF124440.1 gb:NM\_006986.1 gb:AF217963.1

Affy Probe ID	Sequence Description
214875_x_at	Consensus includes gb:AW001847 /FEA=EST /DB_XREF=gi:5848763 /DB_XREF=est:wt81a04.x1 /CLONE=IMAGE:2513838 /UG=Hs.279518 amyloid beta (A4) precursor-like protein 2
211372_s_at	gb:U64094.1 /DEF=Human soluble type II interleukin-1 receptor mRNA, complete cds. /FEA=mRNA /PROD=soluble type II interleukin-1 receptor /DB_XREF=gi:1488065 /UG=Hs.25333 interleukin 1 receptor, type II /FL=gb:U64094.1
205013_s_at	gb:Nm_000675.2 /DEF=Homo sapiens adenosine A2a receptor (ADORA2A), mRNA. /FEA=mRNA /GEN=ADORA2A /PROD=adenosine A2a receptor /DB_XREF=gi:5921991 /UG=Hs.1613 adenosine A2a receptor /FL=gb:M97370.1 gb:Nm_000675.2
212658_at	Consensus includes gb:N66633 /FEA=EST /DB_XREF=gi:1218758 /DB_XREF=est:yy70f02.s1 /CLONE=IMAGE:278907 /UG=Hs.79299 lipoma HMGIC fusion partner-like 2
220161_s_at	gb:Nm_019114.1 /DEF=Homo sapiens EHM2 gene (EHM2), mRNA. /FEA=mRNA /GEN=EHM2 /PROD=EHM2 gene /DB_XREF=gi:9506568 /UG=Hs.267997 EHM2 gene /FL=gb:Nm_019114.1
202194_at	Consensus includes gb:AL117354 /DEF=Human DNA sequence from clone RP5-976O13 on chromosome 1p21.2-22.2 Contains part of the gene for CGI-100 protein, 3 isoforms of the gene for M96 protein, ESTs, STSs, GSSs and a CpG Island /FEA=mRNA_1 /DB_XREF=gi:6822199 /UG=Hs.296155 CGI-100 protein /FL=gb:AF151858.1 gb:Nm_016040.1
213358_at	Consensus includes gb:AB018345.1 /DEF=Homo sapiens mRNA for KIAA0802 protein, partial cds. /FEA=mRNA /GEN=KIAA0802 /PROD=KIAA0802 protein /DB_XREF=gi:3882324 /UG=Hs.27657 KIAA0802 protein
212262_at	Consensus includes gb:AA149639 /FEA=EST /DB_XREF=gi:1720440 /DB_XREF=est:z139c06.s1 /CLONE=IMAGE:504298 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142419.1 gb:AF142422.1
202729_s_at	gb:Nm_000627.1 /DEF=Homo sapiens latent transforming growth factor beta binding protein 1 (LTBP1), mRNA. /FEA=mRNA /GEN=LTBP1 /PROD=latent transforming growth factor beta binding protein 1 precursor /DB_XREF=gi:4557730 /UG=Hs.241257 latent transforming growth factor beta binding protein 1 /FL=gb:M34057.1 gb:Nm_000627.1
211105_s_at	gb:U80918.1 /DEF=Homo sapiens transcription factor (NF-ATcC) mRNA, complete cds. /FEA=mRNA /GEN=NF-ATcC /PROD=transcription factor /DB_XREF=gi:4098860 /UG=Hs.96149 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 /FL=gb:U80918.1
206219_s_at	gb:Nm_005428.2 /DEF=Homo sapiens vav 1 oncogene (VAV1), mRNA. /FEA=mRNA /GEN=VAV1 /PROD=vav 1 oncogene /DB_XREF=gi:7108366 /UG=Hs.116237 vav 1 oncogene /FL=gb:Nm_005428.2



**Affy Probe ID**

**Sequence Description**

210074_at	gb:AF070448.1 /DEF=Homo sapiens cathepsin U (CTSU) mRNA, complete cds. /FEA=mRNA /GEN=CTSU /PROD=cathepsin U /DB_XREF=gi:3228671 /UG=Hs.87417 cathepsin L2 /FL=gb:AB001928.1 gb:AF070448.1 gb:NM_001333.1
203471_s_at	gb:NM_002664.1 /DEF=Homo sapiens plectstrin (PLEK), mRNA. /FEA=mRNA /GEN=PLEK /PROD=pleckstrin /DB_XREF=gi:4505878 /UG=Hs.77436 plectstrin /FL=gb:NM_002664.1
203741_s_at	gb:NM_001114.1 /DEF=Homo sapiens adenylate cyclase 7 (ADCY7), mRNA. /FEA=mRNA /GEN=ADCY7 /PROD=adenylate cyclase 7 /DB_XREF=gi:4557254 /UG=Hs.172199 adenylate cyclase 7 /FL=gb:D25538.1 gb:NM_001114.1
214054_at	Consensus includes gb:AI828929 /FEA=EST /DB_XREF=gi:5449600 /DB_XREF=est:wj37g10.x1 /CLONE=IMAGE:2405058 /UG=Hs.71215 docking protein 2, 56kD /FL=gb:AF034970.1 gb:NM_003974.1
91826_at	Cluster Incl. AI219073:qg16e08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1759718 /clone_end=3 /gb=AI219073 /gi=3801276 /ug=Hs.126062 /len=555
204415_at	gb:NM_022873.1 /DEF=Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 3, mRNA. /FEA=mRNA /GEN=G1P3 /PROD=interferon induced 6-16 protein, isoform c /DB_XREF=gi:13259549 /UG=Hs.265827 interferon, alpha-inducible protein (clone IFI-6-16) /FL=gb:NM_022872.1 gb:NM_022873.1 gb:NM_002038.2
209291_at	Consensus includes gb:AW157094 /FEA=EST /DB_XREF=gi:6228495 /DB_XREF=est:au91g11.x1 /CLONE=IMAGE:2783684 /UG=Hs.34853 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein /FL=gb:NM_001546.1 gb:U16153.1 gb:U28368.1
219560_at	gb:NM_024627.1 /DEF=Homo sapiens hypothetical protein FLJ21125 (FLJ21125), mRNA. /FEA=mRNA /GEN=FLJ21125 /PROD=hypothetical protein FLJ21125 /DB_XREF=gi:13375851 /UG=Hs.36574 hypothetical protein FLJ21125 /FL=gb:NM_024627.1
214521_at	Consensus includes gb:NM_019089.1 /DEF=Homo sapiens hairy and enhancer of split (Drosophila) homolog 2 (HES2), mRNA. /FEA=CDS /GEN=HES2 /PROD=hairy and enhancer of split (Drosophila) homolog2 /DB_XREF=gi:10092644 /UG=Hs.156042 hairy and enhancer of split (Drosophila) homolog 2 /FL=gb:NM_019089.1
202036_s_at	gb:AF017987.1 /DEF=Homo sapiens secreted apoptosis related protein 2 (SARP2) mRNA, complete cds. /FEA=mRNA /GEN=SARP2 /PROD=secreted apoptosis related protein 2 /DB_XREF=gi:2415416 /UG=Hs.7306 secreted frizzled-related protein 1 /FL=gb:AF001900.1 gb:AF017987.1 gb:AF056087.1 gb:NM_003012.2

Affy Probe_ID	Sequence Description
208096_s_at	gb:NM_030820.1 /DEF=Homo sapiens hypothetical protein DKFZp564B052 (DKFZp564B052), mRNA. /FEA=mRNA /GEN=DKFZp564B052 /PROD=hypothetical protein DKFZp564B052 /DB_XREF=gi:13540617 /FL=gb:NM_030820.1
213572_s_at	Consensus includes gb:AI554300 /FEA=EST /DB_XREF=gi:4486663 /DB_XREF=est:10507.x1 /CLONE=IMAGE:2207940 /UG=Hs.183583 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1
219764_at	gb:NM_007197.1 /DEF=Homo sapiens frizzled (Drosophila) homolog 10 (FZD10), mRNA. /FEA=mRNA /GEN=FZD10 /PROD=frizzled (Drosophila) homolog 10 /DB_XREF=gi:6005761 /UG=Hs.31664 frizzled (Drosophila) homolog 10 /FL=gb:AB027464.1 gb:NM_007197.1
218330_s_at	gb:NM_018162.1 /DEF=Homo sapiens hypothetical protein FLJ10633 (FLJ10633), mRNA. /FEA=mRNA /GEN=FLJ10633 /PROD=hypothetical protein FLJ10633 /DB_XREF=gi:9506612 /UG=Hs.23467 hypothetical protein FLJ10633 /FL=gb:NM_018162.1
208616_s_at	gb:U48297.1 /DEF=Homo sapiens protein tyrosine phosphatase PTPCAAX2 (hPTPCAAX2) mRNA, complete cds. /FEA=mRNA /GEN=hPTPCAAX2 /PROD=protein tyrosine phosphatase PTPCAAX2 /DB_XREF=gi:1777756 /UG=Hs.82911 protein tyrosine phosphatase type IVA, member 2 /FL=gb:U48297.1 gb:NM_003479.1 gb:AF208850.1
204754_at	Consensus includes gb:W60800 /FEA=EST /DB_XREF=gi:1367644 /DB_XREF=est:zd27d08.s1 /CLONE=IMAGE:341871 /UG=Hs.250692 hepatic leukemia factor /FL=gb:M95585.1 gb:NM_002126.1
211795_s_at	gb:AF198052.1 /DEF=Homo sapiens EVH1 domain binding protein mRNA, complete cds. /FEA=CDS /PROD=EVH1 domain binding protein /DB_XREF=gi:7416992 /UG=Hs.58435 FYN-binding protein (FYB-120130) /FL=gb:AF198052.1
203927_at	gb:NM_004556.1 /DEF=Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon (NFKBIE), mRNA. /FEA=mRNA /GEN=NFKBIE /PROD=nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon /DB_XREF=gi:4758805 /UG=Hs.91640 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon /FL=gb:U91616.1 gb:NM_004556.1
202917_s_at	gb:NM_002964.2 /DEF=Homo sapiens S100 calcium-binding protein A8 (calgranulin A) (S100A8), mRNA. /FEA=mRNA /GEN=S100A8 /PROD=S100 calcium-binding protein A8 /DB_XREF=gi:9845519 /UG=Hs.100000 S100 calcium-binding protein A8 (calgranulin A) /FL=gb:NM_002964.2
212788_x_at	Consensus includes gb:BG537190 /FEA=EST /DB_XREF=gi:13528922 /DB_XREF=est:602565589F1 /CLONE=IMAGE:4690079 /UG=Hs.111334 ferritin, light polypeptide

Affy Probe_ID	Sequence Description
205606_at	gb:NM_002336.1/DEF=Homo sapiens low density lipoprotein receptor-related protein 6 (LRP6), mRNA. /FEA=mRNA /GEN=LRP6 /PROD=low density lipoprotein receptor-related protein6 /DB_XREF=gi:4505016 /UG=Hs.23672 low density lipoprotein receptor-related protein 6 /FL=gb:AF074264.1 gb:NM_002336.1
209521_s_at	gb:AF286598.1 /DEF=Homo sapiens angiominin mRNA, complete cds. /FEA=mRNA /PROD=angiominin /DB_XREF=gi:9887325 /UG=Hs.9271 KIAA1071 protein /FL=gb:AF286598.1
52255_s_at	Cluster Incl. AI984221.wz57c09.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2562160 /clone_end=3 /gb=AI984221 /gi=5811440 /ug=Hs.235368 /len=515
214056_at	Consensus includes gb:BF981280 /FEA=EST /DB_XREF=gi:12384092 /DB_XREF=est:602308517F1 /CLONE=IMAGE:4399756 /UG=Hs.86386 myeloid cell leukemia sequence 1 (BCL2-related)
36499_at	Cluster Incl. D87469.Human mRNA for KIAA0279 gene, partial cds /cds=(0,7226) /gb=D87469 /gi=1665820 /ug=Hs.57652 /len=8924
203300_x_at	gb:NM_003916.1 /DEF=Homo sapiens adaptor-related protein complex 1, sigma 2 subunit (AP1S2), mRNA. /FEA=mRNA /GEN=AP1S2 /PROD=adaptor-related protein complex 1, sigma 2 subunit /DB_XREF=gi:4506956 /UG=Hs.40368 adaptor-related protein complex 1, sigma 2 subunit /FL=gb:AF251295.1 gb:BC001117.1 gb:AB015320.1 gb:NM_003916.1
205290_s_at	gb:NM_001200.1 /DEF=Homo sapiens bone morphogenetic protein 2 (BMP2), mRNA. /FEA=mRNA /GEN=BMP2 /PROD=bone morphogenetic protein 2 precursor /DB_XREF=gi:4557368 /UG=Hs.73853 bone morphogenetic protein 2 /FL=gb:NM_001200.1
202195_s_at	gb:NM_016040.1 /DEF=Homo sapiens CGI-100 protein (LOC50999), mRNA. /FEA=mRNA /GEN=LOC50999 /PROD=CGI-100 protein /DB_XREF=gi:7705583 /UG=Hs.296155 CGI-100 protein /FL=gb:AF151858.1 gb:NM_016040.1
218125_s_at	gb:NM_018246.1 /DEF=Homo sapiens hypothetical protein FLJ10853 (FLJ10853), mRNA. /FEA=mRNA /GEN=FLJ10853 /PROD=hypothetical protein FLJ10853 /DB_XREF=gi:8922717 /UG=Hs.72085 hypothetical protein FLJ10853 /FL=gb:NM_018246.1
203185_at	gb:NM_014737.1 /DEF=Homo sapiens Ras association (RalGDS/AF-6) domain family 2 (RASSF2), mRNA. /FEA=mRNA /GEN=RASSF2 /PROD=Ras association (RalGDS/AF-6) domain family 2 /DB_XREF=gi:7661963 /UG=Hs.80905 Ras association (RalGDS/AF-6) domain family 2 /FL=gb:D79990.1 gb:NM_014737.1

**Affy Probe ID**

**Sequence Description**

208539\_x\_at gb:NM\_006945.1 /DEF=Homo sapiens small proline-rich protein 2B (SPRR2B), mRNA. /FEA=CDS /GEN=SPRR2B /PROD=small proline-rich protein 2B /DB\_XREF=gi:5902119 /UG=Hs.231622 small proline-rich protein 2B /FL=gb:NM\_006945.1

201852\_x\_at Consensus includes gb:AI813758 /FEA=EST /DB\_XREF=gi:5424973 /DB\_XREF=est:wk79b04.x1 /CLONE=IMAGE:2421583 /UG=Hs.119571 collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant) /FL=gb:NM\_000090.1

204576\_s\_at Consensus includes gb:AA207013 /FEA=EST /DB\_XREF=gi:1802425 /DB\_XREF=est:zr87c08.s1 /CLONE=IMAGE:682670 /UG=Hs.155995 KIAA0643 protein /FL=gb:NM\_024793.1

213256\_at Consensus includes gb:AW593996 /FEA=EST /DB\_XREF=gi:7281254 /DB\_XREF=est:hg41g06.x1 /CLONE=IMAGE:2948218 /UG=Hs.124969 Homo sapiens clone 24707 mRNA sequence

204041\_at gb:NM\_000898.1 /DEF=Homo sapiens monoamine oxidase B (MAOB), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=MAOB /PROD=monoamine oxidase B /DB\_XREF=gi:4505092 /UG=Hs.82163 monoamine oxidase B /FL=gb:M69177.1 gb:NM\_000898.1

202558\_s\_at gb:NM\_006948.1 /DEF=Homo sapiens stress 70 protein chaperone, microsome-associated, 60kD (STCH), mRNA. /FEA=mRNA /GEN=STCH /PROD=stress 70 protein chaperone, microsome-associated, 60kD /DB\_XREF=gi:5902125 /UG=Hs.288799 stress 70 protein chaperone, microsome-associated, 60kD /FL=gb:U04735.1 gb:NM\_006948.1

216891\_at Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CAC)n(GTG)n repeat-containing mRNA. /FEA=mRNA /DB\_XREF=gi:405058 /UG=Hs.104112 Human clone KDB5.1 (CAC)n(GTG)n repeat-containing mRNA

206571\_s\_at gb:NM\_004834.1 /DEF=Homo sapiens mitogen-activated protein kinase kinase kinase 4 (MAP4K4), mRNA. /FEA=mRNA /GEN=MAP4K4 /PROD=mitogen-activated protein kinase kinase kinase 4 /DB\_XREF=gi:4758523 /UG=Hs.3628 mitogen-activated protein kinase kinase kinase 4 /FL=gb:AF096300.1 gb:NM\_004834.1

208992\_s\_at gb:BC000627.1 /DEF=Homo sapiens, Signal transducer and activator of transcription 3, clone MGC:1607, mRNA, complete cds. /FEA=mRNA /PROD=Signal transducer and activator of transcription3 /DB\_XREF=gi:12653684 /UG=Hs.321677 signal transducer and activator of transcription 3 (acute-phase response factor) /FL=gb:BC000627.1 gb:NM\_003150.1 gb:L29277.1

# Affy Probe ID

## Sequence Description

207861\_at gb:NM\_002990.1 /DEF=Homo sapiens small inducible cytokine subfamily A (Cys-Cys), member 22 (SCYA22), mRNA. /FEA=mRNA /GEN=SCYA22 /PROD=small inducible cytokine subfamily A (Cys-Cys), member 22 /DB\_XREF=gi:4506836 /UG=Hs.97203 small inducible cytokine subfamily A (Cys-Cys), member 22 /FL=gb:U83171.1 gb:U83239.1 gb:NM\_002990.1  
218322\_s\_at gb:NM\_016234.2 /DEF=Homo sapiens long-chain fatty acid coenzyme A ligase 5 (FACL5), mRNA. /FEA=mRNA /GEN=FACL5 /PROD=long-chain fatty acid coenzyme A ligase 5 /DB\_XREF=gi:12669912 /UG=Hs.11638 long-chain fatty acid coenzyme A ligase 5 /FL=gb:NM\_016234.2 gb:AB033899.1  
202016\_at gb:NM\_002402.1 /DEF=Homo sapiens mesoderm specific transcript (mouse) homolog (MEST), mRNA. /FEA=mRNA /GEN=MEST /PROD=mesoderm specific transcript (mouse) homolog /DB\_XREF=gi:4505154 /UG=Hs.79284 mesoderm specific transcript (mouse) homolog /FL=gb:BC002413.1 gb:D78611.1 gb:D87367.1 gb:NM\_002402.1 Cluster Incl. AL049370:Homo sapiens mRNA; cDNA DKFZp586D0918 (from clone DKFZp586D0918) /cds=UNKNOWN /gb=AL049370 /gi=4500162 /ug=Hs.13350 /len=2705  
41856\_at Cluster Incl. AF074264:Homo sapiens LDL receptor-related protein 6 (LRP6) mRNA, complete cds /cds=(77,4918) /gb=AF074264 /gi=3462526 /ug=Hs.23672 /len=5288  
34697\_at Consensus includes gb:AI433261 /FEA=EST /DB\_XREF=gi:4287937 /DB\_XREF=est:ti32f08.x1  
222361\_at /CLONE=IMAGE:2132199 /UG=Hs.23189 ESTs, Moderately similar to TBB2\_HUMAN TUBULIN BETA-2 CHAIN H.sapiens  
210335\_at gb:AF056209.1 /DEF=Homo sapiens PAM COOH-terminal interactor protein 1 (PCIP1) mRNA, complete cds. /FEA=mRNA /GEN=PCIP1 /PROD=PAM COOH-terminal interactor protein 1 /DB\_XREF=gi:3560562 /UG=Hs.159396 peptidylglycine alpha-amidating monooxygenase COOH-terminal interactor protein-1 /FL=gb:NM\_005447.2 gb:AF056209.1  
206416\_at gb:NM\_003456.1 /DEF=Homo sapiens zinc finger protein 205 (ZNF205), mRNA. /FEA=mRNA /GEN=ZNF205 /PROD=zinc finger protein 205 /DB\_XREF=gi:4508014 /UG=Hs.13128 zinc finger protein 205 /FL=gb:BC002810.1 gb:AF060865.1 gb:NM\_003456.1  
206400\_at gb:NM\_002307.1 /DEF=Homo sapiens lectin, galactoside-binding, soluble, 7 (galectin 7) (LGALS7), mRNA. /FEA=mRNA /GEN=LGALS7 /PROD=galectin 7 /DB\_XREF=gi:4504984 /UG=Hs.99923 lectin, galactoside-binding, soluble, 7 (galectin 7) /FL=gb:L07769.1 gb:NM\_002307.1 gb:U066643.1

Affy Probe ID	Sequence Description
219696_at	gb:NM_019049.1 /DEF=Homo sapiens hypothetical protein (FLJ20054), mRNA. /FEA=mRNA /GEN=FLJ20054 /PROD=hypothetical protein /DB_XREF=gi:9506654 /UG=Hs.101590 hypothetical protein /FL=gb:NM_019049.1
209366_x_at	gb:M22865.1 /DEF=Human cytochrome b5 mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:181226 /UG=Hs.83834 cytochrome b-5 /FL=gb:M22865.1
208499_s_at	gb:NM_006260.1 /DEF=Homo sapiens protein-kinase, interferon-inducible double stranded RNA dependent inhibitor (PRKRI), mRNA. /FEA=CDS /GEN=PRKRI /PROD=protein-kinase, interferon-inducible doublestranded RNA dependent inhibitor /DB_XREF=gi:5453979 /UG=Hs.9683 DnaJ (Hsp40) homolog, subfamily C, member 3 /FL=gb:U28424.1 gb:NM_006260.1
207076_s_at	gb:NM_000050.1 /DEF=Homo sapiens argininosuccinate synthetase (ASS), mRNA. /FEA=mRNA /GEN=ASS /PROD=argininosuccinate synthetase /DB_XREF=gi:4557336 /UG=Hs.160786 argininosuccinate synthetase /FL=gb:NM_000050.1
209732_at	gb:BC005254.1 /DEF=Homo sapiens, Similar to C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced), clone MGC:12289, mRNA, complete cds. /FEA=mRNA /PROD=Similar to C-type (calcium dependent,carbohydrate-recognition domain) lectin, superfamilymember 2 (activation-induced) /DB_XREF=gi:13528920 /UG=Hs.85201 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced) /FL=gb:BC005254.1 gb:AB015628.1 gb:NM_005127.1
201140_s_at	gb:NM_004583.1 /DEF=Homo sapiens RAB5C, member RAS oncogene family (RAB5C), mRNA. /FEA=mRNA /GEN=RAB5C /PROD=RAB5C, member RAS oncogene family /DB_XREF=gi:4759019 /UG=Hs.479 RAB5C, member RAS oncogene family /FL=gb:NM_004583.1 gb:U11293.1 gb:U18420.1 gb:AF141304.1
212687_at	Consensus includes gb:AL110164.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0324 (from clone DKFZp586I0324). /FEA=mRNA /DB_XREF=gi:5817069 /UG=Hs.193700 Homo sapiens mRNA; cDNA DKFZp586I0324 (from clone DKFZp586I0324)
222303_at	Consensus includes gb:AV700891 /FEA=EST /DB_XREF=gi:10302862 /DB_XREF=est:AV700891 /CLONE=GKCBQD03 /UG=Hs.292477 ESTs
200644_at	gb:NM_023009.1 /DEF=Homo sapiens macrophage myristoylated alanine-rich C kinase substrate (MACMARCKS), mRNA. /FEA=mRNA /GEN=MACMARCKS /PROD=macrophage myristoylated alanine-rich C kinasesubstrate /DB_XREF=gi:13491173 /UG=Hs.75061 macrophage myristoylated alanine-rich C kinase substrate /FL=gb:NM_023009.1

Affy Probe ID	Sequence Description
220528_at	gb:NM_018399.1 /DEF=Homo sapiens VNN3 protein (HSA238982), mRNA. /FEA=mRNA /GEN=HSA238982 /PROD=VNN3 protein /DB_XREF=gi:9055235 /UG=Hs.183656 VNN3 protein /FL=gb:NM_018399.1
209917_s_at	gb:BC002709.1 /DEF=Homo sapiens, TP53 target gene 1, clone MGC:3578, mRNA, complete cds. /FEA=mRNA /PROD=TP53 target gene 1 /DB_XREF=gi:12803742 /UG=Hs.274329 TP53 target gene 1 /FL=gb:BC002709.1
202157_s_at	gb:AB007455.1 gb:NM_007233.1
214582_at	gb:U69546.1 /DEF=Homo sapiens RNA-binding protein BRUNOL3 (BRUNOL3) mRNA, complete cds. /FEA=mRNA /GEN=BRUNOL3 /PROD=RNA-binding protein BRUNOL3 /DB_XREF=gi:1568642 /UG=Hs.211610 CUG triplet repeat, RNA-binding protein 2 /FL=gb:U69546.1 gb:AF036956.1 gb:AF090694.1 gb:NM_006561.1
204589_at	Consensus includes gb:NM_000753.1 /DEF=Homo sapiens 2,3-cyclic nucleotide 3 phosphodiesterase (CNP), mRNA. /FEA=CDS /GEN=CNP /PROD=2,3-cyclic nucleotide 3 phosphodiesterase /DB_XREF=gi:4502924 /UG=Hs.150741 2,3-cyclic nucleotide 3 phosphodiesterase /FL=gb:NM_000753.1
210413_x_at	gb:NM_014840.1 /DEF=Homo sapiens KIAA0537 gene product (KIAA0537), mRNA. /FEA=mRNA /GEN=KIAA0537 /PROD=KIAA0537 gene product /DB_XREF=gi:7662169 /UG=Hs.200598 KIAA0537 gene product /FL=gb:AB011109.1 gb:NM_014840.1
220548_at	gb:U19557.1 /DEF=Human squamous cell carcinoma antigen 2 (SCCA2) mRNA, complete cds. /FEA=mRNA /GEN=SCCA2 /PROD=squamous cell carcinoma antigen 2 /DB_XREF=gi:1052870 /UG=Hs.123035 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 4 /FL=gb:U19557.1
203450_at	gb:NM_006071.1 /DEF=Homo sapiens polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly, sea urchin homolog)-like (PKDREJ), mRNA. /FEA=mRNA /GEN=PKDREJ /PROD=polycystic kidney disease (polycystin) and REJ(sperm receptor for egg jelly, sea urchin homolog)-like /DB_XREF=gi:5174632 /UG=Hs.241383 polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly, sea urchin homolog)-like /FL=gb:AF116458.1 gb:NM_006071.1
205730_s_at	gb:NM_015373.1 /DEF=Homo sapiens chromosome 22 open reading frame 2 (C22ORF2), mRNA. /FEA=mRNA /GEN=C22ORF2 /PROD=chromosome 22 open reading frame 2 /DB_XREF=gi:7656941 /UG=Hs.227637 chromosome 22 open reading frame 2 /FL=gb:AL136686.1 gb:NM_015373.1
	gb:NM_014945.1 /DEF=Homo sapiens KIAA0843 protein (KIAA0843), mRNA. /FEA=mRNA /GEN=KIAA0843 /PROD=KIAA0843 protein /DB_XREF=gi:7662329 /UG=Hs.26777 KIAA0843 protein /FL=gb:AB020650.1
	gb:NM_014945.1

**Affy Probe ID**

**Sequence Description**

218913_s_at	gb:NM_016573.1 /DEF=Homo sapiens Gem-interacting protein (LOC51291), mRNA. /FEA=mRNA /GEN=LOC51291 /PROD=Gem-interacting protein /DB_XREF=gi:7706106 /UG=Hs.49427 Gem-interacting protein /FL=gb:AF132541.1 gb:NM_016573.1
208322_s_at	gb:NM_003033.1 /DEF=Homo sapiens sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase) (SIAT4A), mRNA. /FEA=mRNA /GEN=SIAT4A /PROD=sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase) /DB_XREF=gi:4506950 /UG=Hs.301698 sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase) /FL=gb:L13972.1 gb:NM_003033.1 gb:L29555.1
203473_at	gb:NM_007256.1 /DEF=Homo sapiens solute carrier family 21 (organic anion transporter), member 9 (SLC21A9), mRNA. /FEA=mRNA /GEN=SLC21A9 /PROD=solute carrier family 21 (organic anion transporter), member 9 /DB_XREF=gi:6005819 /UG=Hs.7884 solute carrier family 21 (organic anion transporter), member 9 /FL=gb:AB020687.1 gb:AB026256.1 gb:NM_007256.1
212326_at	Consensus includes gb:AB007922.2 /DEF=Homo sapiens mRNA for KIAA0453 protein, partial cds. /FEA=mRNA /GEN=KIAA0453 /PROD=KIAA0453 protein /DB_XREF=gi:6634036 /UG=Hs.194737 KIAA0453 protein
219760_at	gb:NM_022165.1 /DEF=Homo sapiens Lin-7b protein (LIN-7B), mRNA. /FEA=mRNA /GEN=LIN-7B /PROD=Lin-7b protein /DB_XREF=gi:11545919 /UG=Hs.293686 Lin-7b protein; likely ortholog of mouse LIN-7B; mammalian LIN-7 protein 2 /FL=gb:AF311862.1 gb:NM_022165.1
204321_at	gb:NM_002499.1 /DEF=Homo sapiens neogenin (chicken) homolog 1 (NEO1), mRNA. /FEA=mRNA /GEN=NEO1 /PROD=neogenin (chicken) homolog 1 /DB_XREF=gi:4505374 /UG=Hs.90408 neogenin (chicken) homolog 1 /FL=gb:U61262.1 gb:U72391.1 gb:NM_002499.1
211704_s_at	gb:AF356353.1 /DEF=Homo sapiens spindlin-like protein 2 (SPIN2) mRNA, complete cds. /FEA=mRNA /GEN=SPIN2 /PROD=spindlin-like protein 2 /DB_XREF=gi:13650143 /FL=gb:AF356353.1
205832_at	gb:NM_016352.1 /DEF=Homo sapiens carboxypeptidase A3 (LOC51200), mRNA. /FEA=mRNA /GEN=LOC51200 /PROD=carboxypeptidase A3 /DB_XREF=gi:10047105 /UG=Hs.93764 carboxypeptidase A3 /FL=gb:NM_016352.1 gb:AF095719.1
219694_at	gb:NM_019018.1 /DEF=Homo sapiens hypothetical protein (FLJ11127), mRNA. /FEA=mRNA /GEN=FLJ11127 /PROD=hypothetical protein /DB_XREF=gi:9506640 /UG=Hs.91165 hypothetical protein /FL=gb:NM_019018.1
203733_at	gb:NM_014015.2 /DEF=Homo sapiens MYLE protein (MYLE), mRNA. /FEA=mRNA /GEN=MYLE /PROD=MYLE protein /DB_XREF=gi:13384596 /UG=Hs.11902 MYLE protein /FL=gb:BC001083.1 gb:AF108145.2 gb:NM_014015.2



Affy Probe_ID	Sequence Description
213932_x_at	Consensus includes gb:AI923492 /FEA=EST /DB_XREF=gi:5659456 /DB_XREF=est:wn86a02.x1 /CLONE=IMAGE:2452682 /UG=Hs.181244 major histocompatibility complex, class I, A
203453_at	gb:NM_001038.1 /DEF=Homo sapiens sodium channel, nonvoltage-gated 1 alpha (SCNN1A), mRNA. /FEA=mRNA /GEN=SCNN1A /PROD=sodium channel, nonvoltage-gated 1 alpha /DB_XREF=gi:4506814 /UG=Hs.2794 sodium channel, nonvoltage-gated 1 alpha /FL=gb:NM_001038.1
210164_at	gb:J03189.1 /DEF=Human proteolytic serine esterase-like protein (SECT) gene, complete cds. /FEA=mRNA /DB_XREF=gi:338010 /UG=Hs.1051 granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1) /FL=gb:J04071.1 gb:J03189.1 gb:M17016.1 gb:NM_004131.2
205220_at	gb:NM_006018.1 /DEF=Homo sapiens putative chemokine receptor; GTP-binding protein (HM74), mRNA. /FEA=mRNA /GEN=HM74 /PROD=putative chemokine receptor; GTP-binding protein /DB_XREF=gi:5174460 /UG=Hs.137555 putative chemokine receptor; GTP-binding protein /FL=gb:NM_006018.1
213891_s_at	Consensus includes gb:AI927067 /FEA=EST /DB_XREF=gi:5663031 /DB_XREF=est:wo87f01.x1 /CLONE=IMAGE:2462329 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1000272
204269_at	gb:NM_006875.1 /DEF=Homo sapiens pim-2 oncogene (PIM2), mRNA. /FEA=mRNA /GEN=PIM2 /PROD=pim-2 oncogene /DB_XREF=gi:5803124 /UG=Hs.80205 pim-2 oncogene /FL=gb:U77735.1 gb:NM_006875.1
213050_at	Consensus includes gb:AA594937 /FEA=EST /DB_XREF=gi:2410287 /DB_XREF=est:no40b03.s1 /CLONE=IMAGE:1103117 /UG=Hs.33010 KIAA0633 protein
222293_at	Consensus includes gb:AW204383 /FEA=EST /DB_XREF=gi:6503855 /DB_XREF=est:UI-H-BI1-adv-c-03-0-UI.s1 /CLONE=IMAGE:2717957 /UG=Hs.312723 ESTs, Highly similar to F22162 1 H.sapiens
211725_s_at	gb:BC005884.1 /DEF=Homo sapiens, clone MGC:4736, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:4736) /DB_XREF=gi:13543452 /FL=gb:BC005884.1
204967_at	gb:NM_001649.1 /DEF=Homo sapiens apical protein, Xenopus laevis-like (APXL), mRNA. /FEA=mRNA /GEN=APXL /PROD=apical protein, Xenopus laevis-like /DB_XREF=gi:4502174 /UG=Hs.2391 apical protein, Xenopus laevis-like /FL=gb:NM_001649.1
204284_at	Consensus includes gb:N26005 /FEA=EST /DB_XREF=gi:1140353 /DB_XREF=est:yx88f04.s1 /CLONE=IMAGE:268831 /UG=Hs.303090 protein phosphatase 1, regulatory (inhibitor) subunit 5 /FL=gb:NM_005398.1

**Affy Probe ID**

**Sequence Description**

203127_s_at	gb:BC005123.1 /DEF=Homo sapiens, serine palmitoyltransferase, long chain base subunit 2, clone MGC:10362, mRNA, complete cds. /FEA=mRNA /PROD=serine palmitoyltransferase, long chain basesubunit 2 /DB_XREF=gi:13477298 /UG=Hs.59403 serine palmitoyltransferase, long chain base subunit 2 /FL=gb:BC005123.1 gb:AB011098.1 gb:NM_004863.1
209708_at	gb:AY007239.1 /DEF=Homo sapiens monooxygenase X mRNA, complete cds. /FEA=mRNA /PROD=monooxygenase X /DB_XREF=gi:9988949 /UG=Hs.6909 DKFZP564G202 protein /FL=gb:AY007239.1
210150_s_at	gb:BC003355.1 /DEF=Homo sapiens, clone MGC:5311, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:5311) /DB_XREF=gi:13097167 /UG=Hs.312953 Homo sapiens, clone MGC:5311, mRNA, complete cds /FL=gb:BC003355.1
207480_s_at	gb:NM_020149.1 /DEF=Homo sapiens TALE homeobox protein Meis2e (LOC56908), mRNA. /FEA=mRNA /GEN=LOC56908 /PROD=TALE homeobox protein Meis2e /DB_XREF=gi:9910355 /UG=Hs.283312 TALE homeobox protein Meis2e /FL=gb:AF179899.1 gb:NM_020149.1
202330_s_at	gb:NM_003362.1 /DEF=Homo sapiens uracil-DNA glycosylase (UNG), mRNA. /FEA=mRNA /GEN=UNG /PROD=uracil-DNA glycosylase /DB_XREF=gi:6224978 /UG=Hs.78853 uracil-DNA glycosylase /FL=gb:NM_003362.1
220425_x_at	gb:NM_017578.1 /DEF=Homo sapiens AKAP-binding sperm protein ropporin (DKFZp434B1222), mRNA. /FEA=mRNA /GEN=DKFZp434B1222 /PROD=AKAP-binding sperm protein ropporin /DB_XREF=gi:13487901 /UG=Hs.194093 AKAP-binding sperm protein ropporin /FL=gb:AF231410.1 gb:NM_017578.1
219836_at	gb:NM_024508.1 /DEF=Homo sapiens hypothetical protein MGC10796 (MGC10796), mRNA. /FEA=mRNA /GEN=MGC10796 /PROD=hypothetical protein MGC10796 /DB_XREF=gi:13375643 /UG=Hs.136912 hypothetical protein MGC10796 /FL=gb:BC003536.1 gb:NM_024508.1
220416_at	gb:NM_024837.1 /DEF=Homo sapiens hypothetical protein FLJ21472 (FLJ21472), mRNA. /FEA=mRNA /GEN=FLJ21472 /PROD=hypothetical protein FLJ21472 /DB_XREF=gi:13376248 /UG=Hs.182738 hypothetical protein FLJ21472 /FL=gb:NM_024837.1
210422_x_at	gb:D50402.1 /DEF=Human mRNA for NRAMP1, complete cds. /FEA=mRNA /PROD=Nramp /DB_XREF=gi:1000996 /UG=Hs.182611 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1 /FL=gb:D50402.1 gb:D50403.1 gb:NM_000578.1 gb:L32185.1
213812_s_at	Consensus includes gb:AK024748.1 /DEF=Homo sapiens cDNA: FLJ21095 fis, clone CAS03846. /FEA=mRNA /DB_XREF=gi:10437113 /UG=Hs.108708 calciumcalmodulin-dependent protein kinase 2, beta

Affy Probe_ID	Sequence Description
213392_at	Consensus includes gb:AW070229 /FEA=EST /DB_XREF=gi:6025227 /DB_XREF=est:xa09e01.x1 /CLONE=IMAGE:2567832 /UG=Hs.242407 G protein-coupled receptor, family C, group 5, member B gb:M58596.1 /DEF=Human ELAM-1 ligand fucosyltransferase (ELFT) mRNA, complete cds. /FEA=mRNA /PROD=ELAM-1 ligand fucosyltransferase /DB_XREF=gi:182068 /UG=Hs.2173 fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific) /FL=gb:M58596.1 gb:M58597.1 gb:NM_002033.1
200838_at	gb:NM_001908.1 /DEF=Homo sapiens cathepsin B (CTSB), mRNA. /FEA=mRNA /GEN=CTSB /PROD=cathepsin B /DB_XREF=gi:4503138 /UG=Hs.297939 cathepsin B /FL=gb:M14221.1 gb:L16510.1 gb:NM_001908.1
218018_at	Consensus includes gb:AW449022 /FEA=EST /DB_XREF=gi:6989798 /DB_XREF=est:UI-H-BI3-ake-f-11-0-UI.s1 /CLONE=IMAGE:2734220 /UG=Hs.4746 hypothetical protein FLJ21324 /FL=gb:NM_021941.1 gb:BC003651.1
203724_s_at	gb:NM_014961.1 /DEF=Homo sapiens KIAA0871 protein (KIAA0871), mRNA. /FEA=mRNA /GEN=KIAA0871 /PROD=KIAA0871 protein /DB_XREF=gi:7662351 /UG=Hs.7972 KIAA0871 protein /FL=gb:AB020678.1 gb:NM_014961.1
202789_at	Consensus includes gb:AL022394 /DEF=Human DNA sequence from clone RP3-511B24 on chromosome 20q11.2-12. Contains the 3 end of the TOP1 gene for topoisomerase (DNA) I, the PLOG1 gene for phospholipase C gamma 1, gene KIAA0395 for a possible homeobox protein, a 60S Ribosomal Protein L... /FEA=mRNA_1 /DB_XREF=gi:11345540 /UG=Hs.268177 phospholipase C, gamma 1 (formerly subtype 148) /FL=gb:M34667.1 gb:NM_002660.1
204135_at	gb:NM_014890.1 /DEF=Homo sapiens downregulated in ovarian cancer 1 (DOC1), mRNA. /FEA=mRNA /GEN=DOC1 /PROD=downregulated in ovarian cancer 1 /DB_XREF=gi:7657036 /UG=Hs.15432 downregulated in ovarian cancer 1 /FL=gb:U53445.1 gb:NM_014890.1
211959_at	Consensus includes gb:AW007532 /FEA=EST /DB_XREF=gi:5856310 /DB_XREF=est:ws52h07.x1 /CLONE=IMAGE:2500861 /UG=Hs.103391 Human insulin-like growth factor binding protein 5 (IGFBP5) mRNA gb:NM_024056.1 /DEF=Homo sapiens hypothetical protein MGC5576 (MGC5576), mRNA. /FEA=mRNA
201764_at	/GEN=MGC5576 /PROD=hypothetical protein MGC5576 /DB_XREF=gi:13129025 /UG=Hs.103834 hypothetical protein MGC5576 /FL=gb:BC000854.1 gb:NM_024056.1
200748_s_at	gb:NM_002032.1 /DEF=Homo sapiens ferritin, heavy polypeptide 1 (FTH1), mRNA. /FEA=mRNA /GEN=FTH1 /PROD=ferritin, heavy polypeptide 1 /DB_XREF=gi:4503794 /UG=Hs.62954 ferritin, heavy polypeptide 1 /FL=gb:BC000857.1 gb:BC001399.1 gb:M11146.1 gb:M12937.1 gb:M97164.1 gb:NM_002032.1 gb:L20941.1

Affy Probe_ID	Sequence Description
220049_s_at	gb:NM_025239.1 /DEF=Homo sapiens programmed death ligand 2 (PDL2), mRNA. /FEA=mRNA /GEN=PDL2 /PROD=programmed death ligand 2 /DB_XREF=gi:13376849 /UG=Hs.61929 programmed death ligand 2 /FL=gb:AF344424.1 gb:NM_025239.1
218510_x_at	Consensus includes gb:AI816291 /FEA=EST /DB_XREF=gi:5431837 /DB_XREF=est:au46f08.x1 /CLONE=IMAGE:2517831 /UG=Hs.82273 hypothetical protein /FL=gb:NM_019000.1
206306_at	gb:NM_001036.1 /DEF=Homo sapiens ryanodine receptor 3 (RYR3), mRNA. /FEA=mRNA /GEN=RYR3 /PROD=ryanodine receptor 3 /DB_XREF=gi:4506758 /UG=Hs.9349 ryanodine receptor 3 /FL=gb:AB001025.1 gb:AJ001515.1 gb:NM_001036.1
202935_s_at	Consensus includes gb:AI382146 /FEA=EST /DB_XREF=gi:4194927 /DB_XREF=est:te30c10.x1 /CLONE=IMAGE:2087442 /UG=Hs.2316 SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal) /FL=gb:NM_000346.1
201325_s_at	gb:NM_001423.1 /DEF=Homo sapiens epithelial membrane protein 1 (EMP1), mRNA. /FEA=mRNA /GEN=EMP1 /PROD=epithelial membrane protein 1 /DB_XREF=gi:4503558 /UG=Hs.79368 epithelial membrane protein 1 /FL=gb:U77085.1 gb:U43916.1 gb:NM_001423.1
220856_x_at	gb:NM_014128.1 /DEF=Homo sapiens PRO0470 protein (PRO0470), mRNA. /FEA=mRNA /GEN=PRO0470 /PROD=PRO0470 protein /DB_XREF=gi:7662555 /UG=Hs.278937 PRO0470 protein /FL=gb:AF090928.1 gb:NM_014128.1
209242_at	Consensus includes gb:AL042588 /FEA=EST /DB_XREF=gi:5935496 /DB_XREF=est:DKFZp434K2121_r1 /CLONE=DKFZp434K2121 /UG=Hs.139033 paternally expressed 3 /FL=gb:AF208967.1
219615_s_at	gb:NM_003740.1 /DEF=Homo sapiens potassium channel, subfamily K, member 5 (TASK-2) (KCNK5), mRNA. /FEA=mRNA /GEN=KCNK5 /PROD=potassium channel, subfamily K, member 5 (TASK-2) /DB_XREF=gi:4504850 /UG=Hs.127007 potassium channel, subfamily K, member 5 (TASK-2) /FL=gb:AF084830.1 gb:NM_003740.1
208961_s_at	gb:AB017493.1 /DEF=Homo sapiens mRNA for DNA-binding zinc finger(GBF), complete cds. /FEA=mRNA /PROD=DNA-binding zinc finger(GBF) /DB_XREF=gi:3582142 /UG=Hs.285313 core promoter element binding protein /FL=gb:BC000311.1 gb:BC004301.1 gb:AF001461.1 gb:AB017493.1 gb:NM_001300.2
211382_s_at	gb:AF220152.2 /DEF=Homo sapiens TACC2 mRNA, complete cds. /FEA=mRNA /PROD=TACC2 /DB_XREF=gi:7934571 /UG=Hs.272023 transforming, acidic coiled-coil containing protein 2 /FL=gb:AF220152.2
203640_at	Consensus includes gb:BE328496 /FEA=EST /DB_XREF=gi:9202272 /DB_XREF=est:hs98f09.x1 /CLONE=IMAGE:3145289 /UG=Hs.283609 hypothetical protein PRO2032 /FL=gb:AF116683.1 gb:NM_018615.1

Affy Probe ID	Sequence Description
206453_s_at	gb:NM_016250.1 /DEF=Homo sapiens N-myc downstream-regulated gene 2 (NDRG2), mRNA. /FEA=mRNA /GEN=NDRG2 /PROD=KIAA1248 protein /DB_XREF=gi:10280619 /UG=Hs.243960 N-myc downstream-regulated gene 2 /FL=gb:NM_016250.1 gb:AF159092.3
219066_at	gb:NM_021823.1 /DEF=Homo sapiens hypothetical protein MDS018 (MDS018), mRNA. /FEA=mRNA /GEN=MDS018 /PROD=hypothetical protein MDS018 /DB_XREF=gi:11141896 /UG=Hs.24647 hypothetical protein MDS018 /FL=gb:AF182419.1 gb:NM_021823.1
201328_at	Consensus includes gb:AL575509 /FEA=EST /DB_XREF=gi:12936742 /DB_XREF=est:AL575509 /CLONE=CS0DI059YP21 (3 prime) /UG=Hs.85146 v-ets avian erythroblastosis virus E26 oncogene homolog 2 /FL=gb:J04102.1 gb:NM_005239.1
221655_x_at	gb:AF282167.1 /DEF=Homo sapiens DRC3 mRNA, complete cds. /FEA=mRNA /PROD=DRC3 /DB_XREF=gi:9944939 /UG=Hs.28907 hypothetical protein FLJ20258 /FL=gb:AF282167.1
218779_x_at	gb:NM_017729.1 /DEF=Homo sapiens hypothetical protein FLJ20258 (FLJ20258), mRNA. /FEA=mRNA /GEN=FLJ20258 /PROD=hypothetical protein FLJ20258 /DB_XREF=gi:8923231 /UG=Hs.28907 hypothetical protein FLJ20258 /FL=gb:NM_017729.1
222236_s_at	Consensus includes gb:AK000253.1 /DEF=Homo sapiens cDNA FLJ20246 fis, clone COLF6458. /FEA=mRNA /DB_XREF=gi:7020209 /UG=Hs.44579 hypothetical protein FLJ20199
212071_s_at	Consensus includes gb:BE968833 /FEA=EST /DB_XREF=gi:10579538 /DB_XREF=est:601649861F1 /CLONE=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ13700 fis, clone PLACE2000216, highly similar to SPECTRIN BETA CHAIN, BRAIN
221872_at	Consensus includes gb:AI669229 /FEA=EST /DB_XREF=gi:4834003 /DB_XREF=est:wc13e06.x1 /CLONE=IMAGE:2315074 /UG=Hs.82547 retinoic acid receptor responder (tazarotene induced) 1
209719_x_at	gb:U19556.1 /DEF=Human squamous cell carcinoma antigen 1 (SCCA1) mRNA, complete cds. /FEA=mRNA /GEN=SCCA1 /PROD=squamous cell carcinoma antigen 1 /DB_XREF=gi:1276435 /UG=Hs.227948 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3 /FL=gb:U19556.1 gb:BC005224.1 gb:NM_006919.1
201124_at	Consensus includes gb:AL048423 /FEA=EST /DB_XREF=gi:5936493 /DB_XREF=est:DKFZp586H2324_s1 /CLONE=DKFZp586H2324 /UG=Hs.149846 integrin, beta 5 /FL=gb:M35011.1 gb:J05633.1 gb:NM_002213.1
213455_at	Consensus includes gb:W87466 /FEA=EST /DB_XREF=gi:1401521 /DB_XREF=est:zh67c05.s1 /CLONE=IMAGE:417128 /UG=Hs.246885 hypothetical protein FLJ20783

**Affy Probe\_ID**  
206336\_at

**Sequence Description**

206336\_at

gb:NM\_002993.1 /DEF=Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 6 (granulocyte chemotactic protein 2) (SCYB6), mRNA. /FEA=mRNA /GEN=SCYB6 /PROD=small inducible cytokine subfamily B(Cys-X-Cys), member 6 (granulocyte chemotactic protein 2) /DB\_XREF=gi:4506850 /UG=Hs.164021 small inducible cytokine subfamily B (Cys-X-Cys), member 6 (granulocyte chemotactic protein 2) /FL=gb:U81234.1 gb:NM\_002993.1  
211312\_s\_at gb:AB034725.1 /DEF=Homo sapiens mRNA for WISP-1 variant, complete cds. /FEA=mRNA /PROD=WISP-1 variant /DB\_XREF=gi:11176998 /UG=Hs.194680 WNT1 inducible signaling pathway protein 1 /FL=gb:AB034725.1  
220178\_at gb:NM\_021731.1 /DEF=Homo sapiens hypothetical protein PP3501 (PP3501), mRNA. /FEA=mRNA /GEN=PP3501 /PROD=hypothetical protein PP3501 /DB\_XREF=gi:11119425 /UG=Hs.301406 hypothetical protein PP3501 /FL=gb:NM\_021731.1

205859\_at

gb:NM\_004271.1 /DEF=Homo sapiens MD-1, RP105-associated (MD-1), mRNA. /FEA=mRNA /GEN=MD-1 /PROD=MD-1, RP105-associated /DB\_XREF=gi:4758707 /UG=Hs.184018 MD-1, RP105-associated /FL=gb:AF057178.2 gb:AB020499.1 gb:NM\_004271.1

205464\_at

gb:NM\_000336.1 /DEF=Homo sapiens sodium channel, nonvoltage-gated 1, beta (Liddle syndrome) (SCNN1B), mRNA. /FEA=mRNA /GEN=SCNN1B /PROD=sodium channel, nonvoltage-gated 1, beta /DB\_XREF=gi:4506816 /UG=Hs.37129 sodium channel, nonvoltage-gated 1, beta (Liddle syndrome) /FL=gb:NM\_000336.1 gb:L36593.1  
206388\_at gb:U36798.1 /DEF=Homo sapiens platelet cG1-PDE mRNA, complete cds. /FEA=mRNA /PROD=platelet cG1-PDE /DB\_XREF=gi:1039473 /UG=Hs.777 phosphodiesterase 3A, cGMP-inhibited /FL=gb:U36798.1 gb:M91667.1  
gb:NM\_000921.1

35671\_at

Cluster Incl. U02619:Human TFIIIC Box B-binding subunit mRNA, complete cds /cds=(60,6389) /gb=U02619 /gi=414932 /ug=Hs.331 /len=6996

210793\_s\_at

gb:U41815.1 /DEF=Human nucleoporin 98 (NUP98) mRNA, complete cds. /FEA=mRNA /GEN=NUP98 /PROD=nucleoporin 98 /DB\_XREF=gi:1184172 /UG=Hs.112255 nucleoporin 98kD /FL=gb:U41815.1

208779\_x\_at

gb:L20817.1 /DEF=Homo sapiens tyrosine protein kinase (CAK) gene, complete cds. /FEA=mRNA /GEN=CAK /PROD=tyrosine protein kinase /DB\_XREF=gi:306474 /UG=Hs.75562 discoidin domain receptor family, member 1 /FL=gb:L20817.1

205888\_s\_at

Consensus includes gb:AI962693 /FEA=EST /DB\_XREF=gi:5755406 /DB\_XREF=est:wq55a09.x1 /CLONE=IMAGE:2475160 /UG=Hs.43107 KIAA0555 gene product /FL=gb:AB011127.1 gb:NM\_014790.1

# Affy Probe\_ID

## Sequence Description

220535\_at gb:NM\_018088.1 /DEF=Homo sapiens hypothetical protein FLJ10408 (FLJ10408), mRNA. /FEA=mRNA /GEN=FLJ10408 /PROD=hypothetical protein FLJ10408 /DB\_XREF=gi:8922410 /UG=Hs.196086 hypothetical protein FLJ10408 /FL=gb:NM\_018088.1

206321\_at gb:NM\_002918.1 /DEF=Homo sapiens regulatory factor X, 1 (influences HLA class II expression) (RFX1), mRNA. /FEA=mRNA /GEN=RFX1 /PROD=regulatory factor X, 1 (influences HLA class II expression) /DB\_XREF=gi:4506492 /UG=Hs.123638 regulatory factor X, 1 (influences HLA class II expression) /FL=gb:NM\_002918.1

213293\_s\_at Consensus includes gb:AA083478 /FEA=EST /DB\_XREF=gi:1625564 /DB\_XREF=est:zn32e02.s1 /CLONE=IMAGE:549146 /UG=Hs.295978 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 746039 gb:NM\_015320.1 /DEF=Homo sapiens Rho guanine nucleotide exchange factor (GEF) 4 (ARHGEF4), mRNA. /FEA=mRNA /GEN=ARHGEF4 /PROD=KIAA1112 protein /DB\_XREF=gi:9558722 /UG=Hs.6066 Rho guanine nucleotide exchange factor (GEF) 4 /FL=gb:AF249745.1 gb:NM\_015320.1

205109\_s\_at Consensus includes gb:D50918.1 /DEF=Human mRNA for KIAA0128 gene, partial cds. /FEA=mRNA /GEN=KIAA0128 /DB\_XREF=gi:1469178 /UG=Hs.90998 KIAA0128 protein; septin 2 gb:NM\_018555.2 /DEF=Homo sapiens C2H2-like zinc finger protein (ZNF361), mRNA. /FEA=mRNA /GEN=ZNF361 /PROD=C2H2-like zinc finger protein /DB\_XREF=gi:10092612 /UG=Hs.147644 zinc finger protein 331 /FL=gb:AF251515.2 gb:NM\_018555.2 gb:AF272148.1

212414\_s\_at Consensus includes gb:D50918.1 /DEF=Human mRNA for KIAA0128 gene, partial cds. /FEA=mRNA /GEN=KIAA0128 /DB\_XREF=gi:1469178 /UG=Hs.90998 KIAA0128 protein; septin 2 gb:NM\_018555.2 /DEF=Homo sapiens C2H2-like zinc finger protein (ZNF361), mRNA. /FEA=mRNA /GEN=ZNF361 /PROD=C2H2-like zinc finger protein /DB\_XREF=gi:10092612 /UG=Hs.147644 zinc finger protein 331 /FL=gb:AF251515.2 gb:NM\_018555.2 gb:AF272148.1

220414\_at gb:NM\_017422.2 /DEF=Homo sapiens calmodulin-like skin protein (CLSP), mRNA. /FEA=mRNA /GEN=CLSP /PROD=calmodulin-like skin protein /DB\_XREF=gi:13699870 /UG=Hs.180142 calmodulin-like skin protein /FL=gb:NM\_017422.2 gb:AF172852.1

218548\_x\_at gb:NM\_015926.1 /DEF=Homo sapiens putative secreted protein (SIG11), mRNA. /FEA=mRNA /GEN=SIG11 /PROD=putative secreted protein /DB\_XREF=gi:7706707 /UG=Hs.279785 putative secreted protein /FL=gb:AF072733.1 gb:NM\_015926.1

206170\_at gb:NM\_000024.2 /DEF=Homo sapiens adrenergic, beta-2-, receptor, surface (ADRB2), mRNA. /FEA=mRNA /GEN=ADRB2 /PROD=adrenergic, beta-2-, receptor, surface /DB\_XREF=gi:13162366 /UG=Hs.2551 adrenergic, beta-2-, receptor, surface /FL=gb:NM\_000024.2 gb:M15169.1

Affy Probe_ID	Sequence Description
200800_s_at	gb:NM_005345.3 /DEF=Homo sapiens heat shock 70kD protein 1A (HSPA1A), mRNA. /FEA=mRNA /GEN=HSPA1A /PROD=heat shock 70kD protein 1A /DB_XREF=gi:5579469 /UG=Hs.8997 heat shock 70kD protein 1A /FL=gb:BC002453.1 gb:NM_005345.3
202534_x_at	gb:NM_000791.2 /DEF=Homo sapiens dihydrofolate reductase (DHFR), mRNA. /FEA=mRNA /GEN=DHFR /PROD=dihydrofolate reductase /DB_XREF=gi:7262376 /UG=Hs.83765 dihydrofolate reductase /FL=gb:BC000192.1 gb:BC003584.1 gb:NM_000791.2
202637_s_at	Consensus includes gb:AI608725 /FEA=EST /DB_XREF=gi:4617892 /DB_XREF=est:tw90b01.x1 /CLONE=IMAGE:2266921 /UG=Hs.168383 intercellular adhesion molecule 1 (CD54), human rhinovirus receptor /FL=gb:M24283.1 gb:J03132.1 gb:NM_000201.1
212169_at	Consensus includes gb:AL050187.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1723 (from clone DKFZp586B1723). /FEA=mRNA /DB_XREF=gi:4884402 /UG=Hs.302749 FK506-binding protein 9 (63 kD)
209737_at	gb:AB014605.1 /DEF=Homo sapiens mRNA for KIAA0705 protein, complete cds. /FEA=mRNA /GEN=KIAA0705 /PROD=KIAA0705 protein /DB_XREF=gi:3327223 /UG=Hs.22599 atrophin-1 interacting protein 1; activin receptor interacting protein 1 /FL=gb:AB014605.1
204747_at	gb:NM_001549.1 /DEF=Homo sapiens interferon-induced protein with tetratricopeptide repeats 4 (IFIT4), mRNA. /FEA=mRNA /GEN=IFIT4 /PROD=interferon-induced protein with tetratricopeptide repeats 4 /DB_XREF=gi:4504586 /UG=Hs.181874 interferon-induced protein with tetratricopeptide repeats 4 /FL=gb:BC001383.1 gb:BC004977.1 gb:U52513.1 gb:AF026939.1 gb:AF083470.1 gb:NM_001549.1
222339_x_at	Consensus includes gb:AI054381 /FEA=EST /DB_XREF=gi:3322168 /DB_XREF=est:qi64d09.x1 /CLONE=IMAGE:1861265 /UG=Hs.176139 ESTs
206584_at	gb:NM_015364.1 /DEF=Homo sapiens MD-2 protein (MD-2), mRNA. /FEA=mRNA /GEN=MD-2 /PROD=MD-2 protein /DB_XREF=gi:7662503 /UG=Hs.69328 MD-2 protein /FL=gb:AB018549.1 gb:NM_015364.1 gb:AF168121.1
218469_at	gb:NM_013372.1 /DEF=Homo sapiens cysteine knot superfamily 1, BMP antagonist 1 (CKTTSF1B1), mRNA. /FEA=mRNA /GEN=CKTTSF1B1 /PROD=cysteine knot superfamily 1, BMP antagonist 1 /DB_XREF=gi:7019348 /UG=Hs.40098 cysteine knot superfamily 1, BMP antagonist 1 /FL=gb:AF154054.1 gb:AF045800.1 gb:AF110137.2 gb:NM_013372.1
212188_at	Consensus includes gb:AA551075 /FEA=EST /DB_XREF=gi:2321327 /DB_XREF=est:nk74h06.s1 /CLONE=IMAGE:1019291 /UG=Hs.109438 Homo sapiens clone 24775 mRNA sequence



**Affy Probe\_ID**

**Sequence Description**

209859_at	gb:AF220036.1 /DEF=Homo sapiens tripartite motif protein TRIM9 isoform alpha (TRIM9) mRNA, complete cds; alternatively spliced. /FEA=mRNA /GEN=TRIM9 /PROD=tripartite motif protein TRIM9 isoform alpha /DB_XREF=gi:12407402 /UG=Hs.75090 KIAA0282 protein /FL=gb:AF220036.1 gb:AF220037.1 gb:AF220038.1
211208_s_at	
212158_at	gb:AB039327.2 /DEF=Homo sapiens CASK mRNA for calciumcalmodulin-dependent serine protein kinase, complete cds. /FEA=mRNA /GEN=CASK /PROD=calciumcalmodulin-dependent serine protein kinase /DB_XREF=gi:13359270 /UG=Hs.151469 calciumcalmodulin-dependent serine protein kinase (MAGUK family) /FL=gb:AB039327.2 Consensus includes gb:AL577322 /FEA=EST /DB_XREF=gi:12940338 /DB_XREF=est:AL577322 /CLONE=CS0DI085YI06 (3 prime) /UG=Hs.1501 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
214000_s_at	Consensus includes gb:AI744627 /FEA=EST /DB_XREF=gi:5113004 /DB_XREF=est:wg04g05.x1 /CLONE=IMAGE:2364152 /UG=Hs.82280 regulator of G-protein signalling 10
209583_s_at	gb:AF063591.1 /DEF=Homo sapiens brain my033 protein mRNA, complete cds. /FEA=mRNA /PROD=brain my033 protein /DB_XREF=gi:12002013 /UG=Hs.79015 antigen identified by monoclonal antibody MRC OX-2 /FL=gb:AF063591.1
214255_at	Consensus includes gb:N35112 /FEA=EST /DB_XREF=gi:1156254 /DB_XREF=est:yy19h05.s1 /CLONE=IMAGE:271737 /UG=Hs.44697 ATPase, Class V, type 10C
221828_s_at	Consensus includes gb:AK024432.1 /DEF=Homo sapiens mRNA for FLJ00022 protein, partial cds. /FEA=mRNA /GEN=FLJ00022 /PROD=FLJ00022 protein /DB_XREF=gi:10440372 /UG=Hs.4914 Homo sapiens mRNA for FLJ00031 protein, partial cds
203315_at	gb:BC000103.1 /DEF=Homo sapiens, NCK adaptor protein 2, clone MGC:1698, mRNA, complete cds. /FEA=mRNA /PROD=NCK adaptor protein 2 /DB_XREF=gi:12652708 /UG=Hs.101695 NCK adaptor protein 2 /FL=gb:BC000103.1 gb:AF043119.1 gb:AF047487.1 gb:NM_003581.1
219861_at	gb:NM_018163.1 /DEF=Homo sapiens hypothetical protein FLJ10634 (FLJ10634), mRNA. /FEA=mRNA /GEN=FLJ10634 /PROD=hypothetical protein FLJ10634 /DB_XREF=gi:8922562 /UG=Hs.24989 hypothetical protein FLJ10634 /FL=gb:BC000048.1 gb:NM_018163.1

Affy Probe_ID	Sequence Description
213540_at	Consensus includes gb:AL031228 /DEF=Human DNA sequence from clone 1033B10 on chromosome 6p21.2-21.31. Contains the BING5 gene, exons 11 to 15 of the BING4 gene, the gene for GalT3 (beta3-Galactosyltransferase), the RPS18 (40S ribosomal protein S18) gene, the SACM2L (suppressor of ac... /FEA=mRNA_1 /DB_XREF=gi:3646023 /UG=Hs.288354 FabG (beta-ketoacyl-acyl-carrier-protein reductase, E coli) like gb:NM_030763.1 /DEF=Homo sapiens nucleosomal binding protein 1 (NSBP1), mRNA. /FEA=mRNA /GEN=NSBP1 /PROD=nucleosomal binding protein 1 /DB_XREF=gi:13540522 /FL=gb:NM_030763.1 Consensus includes gb:H49382 /FEA=EST /DB_XREF=gi:989223 /DB_XREF=est:yq19b06.s1 /CLONE=IMAGE:274115 /UG=Hs.124984 ESTs, Moderately similar to ALU7_HUMAN ALU SUBFAMILY SQ SEQUENCE CONTAMINATION WARNING ENTRY H.sapiens
221238_at	gb:NM_021158.1 /DEF=Homo sapiens protein kinase domains containing protein similar to phosphoprotein C8FW (LOC57761), mRNA. /FEA=mRNA /GEN=LOC57761 /PROD=protein kinase domains containing proteins similar to phosphoprotein C8FW /DB_XREF=gi:11056039 /UG=Hs.26802 protein kinase domains containing protein similar to phosphoprotein C8FW /FL=gb:NM_021158.1
214051_at	Consensus includes gb:BF677486 /FEA=EST /DB_XREF=gi:11951381 /DB_XREF=est:602085745F1 /CLONE=IMAGE:4249834 /UG=Hs.56145 thymosin, beta, identified in neuroblastoma cells
211776_s_at	gb:BC006141.1 /DEF=Homo sapiens, Similar to erythrocyte protein band 4.1-like 3, clone MGC:13087, mRNA, complete cds. /FEA=mRNA /PROD=Similar to erythrocyte protein band 4.1-like 3 /DB_XREF=gi:13544008 /FL=gb:BC006141.1
203874_s_at	gb:NM_003069.1 /DEF=Homo sapiens SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 (SMARCA1), mRNA. /FEA=mRNA /GEN=SMARCA1 /PROD=SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 /DB_XREF=gi:4507066 /UG=Hs.152292 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 /FL=gb:M88163.1 gb:NM_003069.1
202552_s_at	gb:NM_016441.1 /DEF=Homo sapiens cysteine-rich motor neuron 1 (CRIM1), mRNA. /FEA=mRNA /GEN=CRIM1 /PROD=cysteine-rich motor neuron 1 /DB_XREF=gi:10092638 /UG=Hs.19280 cysteine-rich motor neuron 1 /FL=gb:NM_016441.1 gb:AF167706.1
213578_at	Consensus includes gb:AI678679 /FEA=EST /DB_XREF=gi:4888861 /DB_XREF=est:tu58f02.x1 /CLONE=IMAGE:2255259 /UG=Hs.2534 bone morphogenetic protein receptor, type IA

**Affy Probe\_ID**

**Sequence Description**

204362\_at

gb:NM\_003930.1 /DEF=Homo sapiens SKAP55 homologue (SKAP-HOM), mRNA. /FEA=mRNA /GEN=SKAP-HOM /PROD=SKAP55 homologue /DB\_XREF=gi:4506962 /UG=Hs.52644 SKAP55 homologue /FL=gb:BC002893.1 gb:AF072166.1 gb:AB014486.1 gb:AF051323.1 gb:NM\_003930.1

209094\_at

Consensus includes gb:AL078459 /DEF=Human DNA sequence from clone RP4-621F18 on chromosome 1p11.4-21.3. Contains the 3 end of the gene for ng.ng dimethylarginine dimethylaminohydrolase (EC 3.5.3.18), ESTs, STSs and GSSs /FEA=mRNA /DB\_XREF=gi:5791502 /UG=Hs.303180 dimethylarginine dimethylaminohydrolase 1 /FL=gb:AB001915.1 gb:NM\_012137.1

219477\_s\_at

gb:NM\_018676.1 /DEF=Homo sapiens TMTSP for transmembrane molecule with thrombospondin module (LOC55901), mRNA. /FEA=mRNA /GEN=LOC55901 /PROD=TMTSP for transmembrane molecule with thrombospondin module /DB\_XREF=gi:8923893 /UG=Hs.325667 TMTSP for transmembrane molecule with thrombospondin module /FL=gb:AB044385.1 gb:NM\_018676.1

218035\_s\_at

gb:NM\_019027.1 /DEF=Homo sapiens hypothetical protein (FLJ20273), mRNA. /FEA=mRNA /GEN=FLJ20273 /PROD=hypothetical protein /DB\_XREF=gi:9506670 /UG=Hs.95549 hypothetical protein /FL=gb:NM\_019027.1 Consensus includes gb:BG251266 /FEA=EST /DB\_XREF=gi:12761082 /DB\_XREF=est:602365892F1 /CLONE=IMAGE:4474055 /UG=Hs.283565 FOS-like antigen-1 /FL=gb:NM\_005438.1

204420\_at

Consensus includes gb:BG230614 /FEA=EST /DB\_XREF=gi:12725656 /DB\_XREF=est:naf41b12.x1 Consensus includes gb:4143335 /UG=Hs.82685 CD47 antigen (Rh-related antigen, integrin-associated signal transducer) /CLONE=IMAGE:4143335 /UG=Hs.82685 CD47 antigen (Rh-related antigen, integrin-associated signal transducer) gb:AF091627.1 /DEF=Homo sapiens CUSP mRNA, complete cds. /FEA=mRNA /PROD=CUSP /DB\_XREF=gi:3644039 /UG=Hs.137569 tumor protein 63 kDa with strong homology to p53 /FL=gb:AF091627.1 gb:AF075431.1 gb:AF075433.1

213857\_s\_at

gb:NM\_002041.2 /DEF=Homo sapiens GA-binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA. /FEA=mRNA /GEN=GABPB2 /PROD=GA-binding protein transcription factor, betasubunit 2 (47kD), isoform gamma 1 /DB\_XREF=gi:8051596 /UG=Hs.278238 GA-binding protein transcription factor, beta subunit 2 (47kD) /FL=gb:U13047.1 gb:NM\_002041.2

209863\_s\_at

gb:NM\_016951.2 /DEF=Homo sapiens transmembrane proteolipid (HSPC224), mRNA. /FEA=mRNA /GEN=HSPC224 /PROD=transmembrane proteolipid /DB\_XREF=gi:10092593 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM\_016951.2 gb:BC004380.1 gb:AF057306.1 gb:AF151058.1 gb:AF135380.2

206173\_x\_at

gb:NM\_002041.2 /DEF=Homo sapiens GA-binding protein transcription factor, beta subunit 2 (47kD) (GABPB2), transcript variant gamma, mRNA. /FEA=mRNA /GEN=GABPB2 /PROD=GA-binding protein transcription factor, betasubunit 2 (47kD), isoform gamma 1 /DB\_XREF=gi:8051596 /UG=Hs.278238 GA-binding protein transcription factor, beta subunit 2 (47kD) /FL=gb:U13047.1 gb:NM\_002041.2

219161\_s\_at

gb:NM\_016951.2 /DEF=Homo sapiens transmembrane proteolipid (HSPC224), mRNA. /FEA=mRNA /GEN=HSPC224 /PROD=transmembrane proteolipid /DB\_XREF=gi:10092593 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM\_016951.2 gb:BC004380.1 gb:AF057306.1 gb:AF151058.1 gb:AF135380.2

# Affy Probe ID

## Sequence Description

220146\_at gb:NM\_016562.1 /DEF=Homo sapiens toll-like receptor 7 (LOC51284), mRNA. /FEA=mRNA /GEN=LOC51284 /PROD=toll-like receptor 7 /DB\_XREF=gi:7706092 /UG=Hs.179152 toll-like receptor 7 /FL=gb:AF240467.1 gb:NM\_016562.1 gb:AF245702.1  
202266\_at gb:NM\_016614.1 /DEF=Homo sapiens TRAF and TNF receptor-associated protein (AD022), mRNA. /FEA=mRNA /GEN=AD022 /PROD=TRAF and TNF receptor-associated protein /DB\_XREF=gi:7705261 /UG=Hs.46847 TRAF and TNF receptor-associated protein /FL=gb:AF201687.1 gb:AF223469.1 gb:NM\_016614.1  
204834\_at gb:NM\_006682.1 /DEF=Homo sapiens fibrinogen-like 2 (FGL2), mRNA. /FEA=mRNA /GEN=FGL2 /PROD=fibrinogen-like 2 /DB\_XREF=gi:5730074 /UG=Hs.2659 fibrinogen-like 2 /FL=gb:NM\_006682.1  
204401\_at gb:NM\_002250.1 /DEF=Homo sapiens potassium intermediate small conductance calcium-activated channel, subfamily N, member 4 (KCNN4), mRNA. /FEA=mRNA /GEN=KCNN4 /PROD=potassium intermediate small conductance calcium-activated channel, subfamily N, member 4 /DB\_XREF=gi:4504858 /UG=Hs.10082 potassium intermediate small conductance calcium-activated channel, subfamily N, member 4 /FL=gb:AF000972.1 gb:AF033021.1 gb:AF022150.1 gb:AF022797.1 gb:NM\_002250.1  
220894\_x\_at gb:NM\_021619.1 /DEF=Homo sapiens PR domain containing 12 (PRDM12), mRNA. /FEA=mRNA /GEN=PRDM12 /PROD=PR domain containing 12 /DB\_XREF=gi:11055979 /UG=Hs.287385 PR domain containing 12 /FL=gb:AY004252.1 gb:NM\_021619.1  
215075\_s\_at Consensus includes gb:L29511.1 /DEF=Human GRB2 isoform mRNA. /FEA=mRNA /PROD=growth factor receptor-bound protein 3 /DB\_XREF=gi:460667 /UG=Hs.296381 growth factor receptor-bound protein 2  
201983\_s\_at Consensus includes gb:AW157070 /FEA=EST /DB\_XREF=gi:6228471 /DB\_XREF=est:au91e07.x1 /CLONE=IMAGE:2783652 /UG=Hs.77432 epidermal growth factor receptor (avian erythroblastic leukemia viral (v-erb-b) oncogene homolog) /FL=gb:NM\_005228.1  
208436\_s\_at gb:NM\_004030.1 /DEF=Homo sapiens interferon regulatory factor 7 (IRF7), transcript variant c, mRNA. /FEA=mRNA /GEN=IRF7 /PROD=interferon regulatory factor 7, transcript variant c /DB\_XREF=gi:4809285 /UG=Hs.166120 interferon regulatory factor 7 /FL=gb:NM\_004030.1  
219622\_at gb:NM\_017817.1 /DEF=Homo sapiens hypothetical protein FLJ20429 (FLJ20429), mRNA. /FEA=mRNA /GEN=FLJ20429 /PROD=hypothetical protein FLJ20429 /DB\_XREF=gi:89233400 /UG=Hs.179791 hypothetical protein FLJ20429 /FL=gb:NM\_017817.1

Affy Probe_ID	Sequence Description
207232_s_at	gb:NM_014648.1 /DEF=Homo sapiens KIAA0675 gene product (KIAA0675), mRNA. /FEA=mRNA /GEN=KIAA0675 /PROD=KIAA0675 gene product /DB_XREF=gi:7662243 /UG=Hs.165662 KIAA0675 gene product /FL=gb:AB014575.1 gb:NM_014648.1
203105_s_at	gb:NM_012062.1 /DEF=Homo sapiens dynamin 1-like (DNM1L), transcript variant 1, mRNA. /FEA=mRNA /GEN=DNM1L /PROD=dynamin 1-like protein, isoform 1 /DB_XREF=gi:6996004 /UG=Hs.180628 dynamin 1-like /FL=gb:AF000430.1 gb:AB006965.1 gb:NM_012062.1
215440_s_at	Consensus includes gb:AL523320 /FEA=EST /DB_XREF=gi:12786813 /DB_XREF=est:AL523320 /CLONE=CS0DC001YD02 (5 prime) /UG=Hs.184736 hypothetical protein FLJ10097
203778_at	gb:NM_005908.1 /DEF=Homo sapiens mannosidase, beta A, lysosomal (MANBA), mRNA. /FEA=mRNA /GEN=MANBA /PROD=mannosidase, beta A, lysosomal /DB_XREF=gi:5174522 /UG=Hs.115945 mannosidase, beta A, lysosomal /FL=gb:U60337.1 gb:NM_005908.1
204614_at	gb:NM_002575.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 (SERPINB2), mRNA. /FEA=mRNA /GEN=SERPINB2 /PROD=serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 /DB_XREF=gi:4505594 /UG=Hs.75716 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2 /FL=gb:J02685.1 gb:J03603.1 gb:M18082.1 gb:NM_002575.1
209590_at	Consensus includes gb:AL157414 /DEF=Human DNA sequence from clone RP11-560A15 on chromosome 20 Contains part of a novel gene, the 3 part of the BMP7 (bone morphogenetic protein 7 (osteogenic protein 1)) gene, ESTs, STSs, GSSs and a CpG island /FEA=mRNA_1 /DB_XREF=gi:9581576 /UG=Hs.170195 bone morphogenetic protein 7 (osteogenic protein 1) /FL=gb:M60316.1 gb:NM_001719.1
204621_s_at	Consensus includes gb:AI935096 /FEA=EST /DB_XREF=gi:5673966 /DB_XREF=est:wp13e10.x1 /CLONE=IMAGE:2464746 /UG=Hs.82120 nuclear receptor subfamily 4, group A, member 2 /FL=gb:NM_006186.1
204998_s_at	gb:NM_012068.2 /DEF=Homo sapiens activating transcription factor 5 (ATF5), mRNA. /FEA=mRNA /GEN=ATF5 /PROD=activating transcription factor 5 /DB_XREF=gi:12597624 /UG=Hs.9754 activating transcription factor 5 /FL=gb:AF305687.1 gb:AB021663.2 gb:NM_012068.2 gb:BC005174.1
204881_s_at	gb:NM_003358.1 /DEF=Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA. /FEA=mRNA /GEN=UGCG /PROD=ceramide glucosyltransferase /DB_XREF=gi:4507810 /UG=Hs.152601 UDP-glucose ceramide glucosyltransferase /FL=gb:D50840.1 gb:NM_003358.1
37892_at	Cluster Incl. J04177:Human alpha-1 type XI collagen (COL11A1) mRNA, complete cds /cds=(161,5581) /gb=J04177 /gi=179729 /ug=Hs.82772 /len=6158

# Affy Probe\_ID

## Sequence Description

201669\_s\_at gb:NM\_002356.4 /DEF=Homo sapiens myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) (MACS), mRNA. /FEA=mRNA /GEN=MACS /PROD=myristoylated alanine-rich protein kinase C substrate /DB\_XREF=gi:11125771 /UG=Hs.75607 myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L) /FL=gb:NM\_002356.4 gb:M68956.1 gb:D10522.1

219517\_at gb:NM\_025165.1 /DEF=Homo sapiens hypothetical protein FLJ22637 (FLJ22637), mRNA. /FEA=mRNA /GEN=FLJ22637 /PROD=hypothetical protein FLJ22637 /DB\_XREF=gi:13376767 /UG=Hs.296178 hypothetical protein FLJ22637 /FL=gb:NM\_025165.1

202562\_s\_at gb:AL136658.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564H0664 (from clone DKFZp564H0664); complete cds. /FEA=mRNA /GEN=DKFZp564H0664 /PROD=hypothetical protein /DB\_XREF=gi:12052839 /UG=Hs.15106 chromosome 14 open reading frame 1 /FL=gb:AL136658.1 gb:AF136971.1 gb:BC002444.1 gb:AF134159.2 gb:NM\_007176.1

201416\_at Consensus includes gb:BG528420 /FEA=EST /DB\_XREF=gi:13519957 /DB\_XREF=est:602579853F1 /CLONE=IMAGE:4719060 /UG=Hs.83484 SRY (sex determining region Y)-box 4 /FL=gb:NM\_003107.1

219210\_s\_at gb:NM\_016530.1 /DEF=Homo sapiens RAB-8b protein (LOC51762), mRNA. /FEA=mRNA /GEN=LOC51762 /PROD=RAB-8b protein /DB\_XREF=gi:7706562 /UG=Hs.321245 RAB-8b protein /FL=gb:AB038995.1 gb:NM\_016530.1

208875\_s\_at Consensus includes gb:BF796470 /FEA=EST /DB\_XREF=gi:12101524 /DB\_XREF=est:602259926F1 /CLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1

202073\_at Consensus includes gb:AV757675 /FEA=EST /DB\_XREF=gi:10915523 /DB\_XREF=est:AV757675 /CLONE=BMFAVB12 /UG=Hs.278898 tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein /FL=gb:NM\_021980.1

221991\_at Consensus includes gb:AI937333 /FEA=EST /DB\_XREF=gi:5676203 /DB\_XREF=est:wp75h10.x1 /CLONE=IMAGE:2467651 /UG=Hs.55069 neurexophilin 3

201847\_at gb:NM\_000235.1 /DEF=Homo sapiens lipase A, lysosomal acid, cholesterol esterase (Wolman disease) (LIPA), mRNA. /FEA=mRNA /GEN=LIPA /PROD=lipase A precursor /DB\_XREF=gi:4557720 /UG=Hs.85226 lipase A, lysosomal acid, cholesterol esterase (Wolman disease) /FL=gb:M74775.1 gb:NM\_000235.1 gb:U08464.1

Affy Probe ID	Sequence Description
203178_at	gb:NM_001482.1 /DEF=Homo sapiens glycine amidinotransferase (L-arginine:glycine amidinotransferase) (GATM), mRNA. /FEA=mRNA /GEN=GATM /PROD=glycine amidinotransferase (L-arginine:glycine amidinotransferase) /DB_XREF=gi:4503932 /UG=Hs.75335 glycine amidinotransferase (L-arginine:glycine amidinotransferase) /FL=gb:BC004141.1 gb:NM_001482.1
221009_s_at	gb:NM_016109.1 /DEF=Homo sapiens PPAR(gamma) angiopoietin related protein (PGAR), mRNA. /FEA=mRNA /GEN=PGAR /PROD=PPAR(gamma) angiopoietin related protein /DB_XREF=gi:7705828 /UG=Hs.9613 PPAR(gamma) angiopoietin related protein /FL=gb:AF153606.1 gb:NM_016109.1
216935_at	Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=gi:2589191 /UG=Hs.113261 Homo sapiens skin-specific protein (xp33) mRNA, partial cds
210041_s_at	gb:BC001258.1 /DEF=Homo sapiens, N-acetylglucosamine-phosphate mutase, clone MGC:5002, mRNA, complete cds. /FEA=mRNA /PROD=N-acetylglucosamine-phosphate mutase /DB_XREF=gi:12654830 /UG=Hs.237323 N-acetylglucosamine-phosphate mutase /FL=gb:BC001258.1 gb:AF102265.1 gb:AF180371.1 gb:AB032081.1
218788_s_at	gb:NM_022743.1 /DEF=Homo sapiens hypothetical protein FLJ21080 (FLJ21080), mRNA. /FEA=mRNA /GEN=FLJ21080 /PROD=hypothetical protein FLJ21080 /DB_XREF=gi:12232400 /UG=Hs.8109 hypothetical protein FLJ21080 /FL=gb:NM_022743.1
202647_s_at	gb:NM_002524.2 /DEF=Homo sapiens neuroblastoma RAS viral (v-ras) oncogene homolog (NRAS), mRNA. /FEA=mRNA /GEN=NRAS /PROD=neuroblastoma RAS viral (v-ras) oncogene homolog /DB_XREF=gi:6006027 /UG=Hs.260523 neuroblastoma RAS viral (v-ras) oncogene homolog /FL=gb:BC005219.1 gb:NM_002524.2
217585_at	Consensus includes gb:BE502910 /FEA=EST /DB_XREF=gi:9705318 /DB_XREF=est:h281a09.x1
207719_x_at	/CLONE=IMAGE:3214360 /UG=Hs.166449 ESTs, Weakly similar to S55024 nebulin, skeletal muscle H.sapiens gb:NM_014812.1 /DEF=Homo sapiens KIAA0470 gene product (KIAA0470), mRNA. /FEA=mRNA /GEN=KIAA0470 /PROD=KARP-1-binding protein /DB_XREF=gi:7662141 /UG=Hs.25132 KIAA0470 gene product /FL=gb:AB022657.1 gb:NM_014812.1
207038_at	gb:NM_004694.1 /DEF=Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 6 (SLC16A6), mRNA. /FEA=mRNA /GEN=SLC16A6 /PROD=solute carrier family 16 (monocarboxylic acid transporters), member 6 /DB_XREF=gi:4759117 /UG=Hs.114924 solute carrier family 16 (monocarboxylic acid transporters), member 6 /FL=gb:U79745.1 gb:NM_004694.1

# Affy Probe\_ID

## Sequence Description

201904\_s\_at Consensus includes gb:BF031714 /FEA=EST /DB\_XREF=gi:10739426 /DB\_XREF=est:601558023F1 /CLONE=IMAGE:3827909 /UG=Hs.147189 HYA22 protein /FL=gb:D88153.1 gb:NM\_005808.1

219076\_s\_at gb:NM\_018663.1 /DEF=Homo sapiens 22kDa peroxisomal membrane protein-like (LOC55895), mRNA. /FEA=mRNA /GEN=LOC55895 /PROD=22kDa peroxisomal membrane protein-like /DB\_XREF=gi:8923891 /UG=Hs.49912 22kDa peroxisomal membrane protein-like /FL=gb:AF250136.1 gb:NM\_018663.1

221133\_s\_at gb:NM\_016369.1 /DEF=Homo sapiens claudin 18 (CLDN18), mRNA. /FEA=mRNA /GEN=CLDN18 /PROD=claudin 18 /DB\_XREF=gi:7705960 /UG=Hs.278966 Claudin-18 /FL=gb:AF221069.1 gb:NM\_016369.1

203693\_s\_at gb:NM\_001949.2 /DEF=Homo sapiens E2F transcription factor 3 (E2F3) mRNA, complete cds. /FEA=mRNA /GEN=E2F3 /PROD=E2F transcription factor 3 /DB\_XREF=gi:12669913 /UG=Hs.1189 E2F transcription factor 3 /FL=gb:NM\_001949.2

212077\_at Consensus includes gb:AL583520 /FEA=EST /DB\_XREF=gi:12952562 /DB\_XREF=est:AL583520 /CLONE=CS0DC024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mRNA for caldesmon, 3 UTR

212297\_at Consensus includes gb:BF218804 /FEA=EST /DB\_XREF=gi:11112494 /DB\_XREF=est:601882315F1 /CLONE=IMAGE:4094786 /UG=Hs.4934 H.sapiens polyA site DNA

219587\_at gb:NM\_017868.1 /DEF=Homo sapiens hypothetical protein FLJ20535 (FLJ20535), mRNA. /FEA=mRNA /GEN=FLJ20535 /PROD=hypothetical protein FLJ20535 /DB\_XREF=gi:8923504 /UG=Hs.183362 hypothetical protein FLJ20535 /FL=gb:NM\_017868.1

218610\_s\_at gb:NM\_018340.1 /DEF=Homo sapiens hypothetical protein FLJ11151 (FLJ11151), mRNA. /FEA=mRNA /GEN=FLJ11151 /PROD=hypothetical protein FLJ11151 /DB\_XREF=gi:8922900 /UG=Hs.14992 hypothetical protein FLJ11151 /FL=gb:NM\_018340.1

200889\_s\_at Consensus includes gb:AI016620 /FEA=EST /DB\_XREF=gi:3230956 /DB\_XREF=est:ov30e10.x1 /CLONE=IMAGE:1638858 /UG=Hs.250773 signal sequence receptor, alpha (translocon-associated protein alpha) /FL=gb:AF156965.1 gb:NM\_003144.2

204319\_s\_at gb:NM\_002925.2 /DEF=Homo sapiens regulator of G-protein signalling 10 (RGS10), mRNA. /FEA=mRNA /GEN=RGS10 /PROD=regulator of G-protein signaling 10 /DB\_XREF=gi:1184225 /UG=Hs.82280 regulator of G-protein signalling 10 /FL=gb:NM\_002925.2 gb:AF045229.1

205531\_s\_at gb:NM\_013267.1 /DEF=Homo sapiens breast cell glutaminase (GA), mRNA. /FEA=mRNA /GEN=GA /PROD=breast cell glutaminase /DB\_XREF=gi:7019388 /UG=Hs.325443 breast cell glutaminase /FL=gb:AF110330.1 gb:AF110331.1 gb:AF223944.1 gb:NM\_013267.1



# Affy Probe\_ID

## Sequence Description

213413\_at Consensus includes gb:BG434174 /FEA=EST /DB\_XREF=gi:13340680 /DB\_XREF=est:602506106F1 /CLONE=IMAGE:4603513 /UG=Hs.23437 Homo sapiens cDNA FLJ13555 fis, clone PLACE1007677 gb:NM\_024944.1 /DEF=Homo sapiens hypothetical protein FLJ12627 (FLJ12627), mRNA. /FEA=mRNA /GEN=FLJ12627 /PROD=hypothetical protein FLJ12627 /DB\_XREF=gi:13376424 /UG=Hs.283725 hypothetical protein FLJ12627 /FL=gb:NM\_024944.1

202422\_s\_at gb:NM\_022977.1 /DEF=Homo sapiens fatty-acid-Coenzyme A ligase, long-chain 4 (FACL4), transcript variant 2, mRNA. /FEA=mRNA /GEN=FACL4 /PROD=long-chain fatty-acid-Coenzyme A ligase 4, isoform 2 /DB\_XREF=gi:12669908 /UG=Hs.81452 fatty-acid-Coenzyme A ligase, long-chain 4 /FL=gb:NM\_022977.1

216640\_s\_at Consensus includes gb:AK026926.1 /DEF=Homo sapiens cDNA: FLJ23273 fis, clone HEP02611, highly similar to HSU79278 Human protein disulfide isomerase-related protein P5 mRNA. /FEA=mRNA /DB\_XREF=gi:10439898 /UG=Hs.182429 protein disulfide isomerase-related protein

201286\_at Consensus includes gb:Z48199 /DEF=H.sapiens syndecan-1 gene (exons 2-5) /FEA=mRNA /DB\_XREF=gi:666051 /UG=Hs.82109 syndecan 1 /FL=gb:J05392.1 gb:NM\_002997.1

36030\_at Cluster Incl. AL080214:Homo sapiens mRNA; cDNA DKFZp586I2223 (from clone DKFZp586I2223) /cds=(0,318) /gb=AL080214 /gi=5262704 /ug=Hs.46659 /len=1272

202936\_s\_at gb:NM\_000346.1 /DEF=Homo sapiens SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal) (SOX9), mRNA. /FEA=mRNA /GEN=SOX9 /PROD=transcription factor SOX9 /DB\_XREF=gi:4557852 /UG=Hs.2316 SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal) /FL=gb:NM\_000346.1

206924\_at gb:NM\_000641.1 /DEF=Homo sapiens interleukin 11 (IL11), mRNA. /FEA=mRNA /GEN=IL11 /PROD=interleukin 11 /DB\_XREF=gi:10834993 /UG=Hs.1721 interleukin 11 /FL=gb:NM\_000641.1 gb:M57765.1

204454\_at gb:NM\_012317.1 /DEF=Homo sapiens leucine zipper, down-regulated in cancer 1 (LDOC1), mRNA. /FEA=mRNA /GEN=LDOC1 /PROD=leucine zipper, down-regulated in cancer 1 /DB\_XREF=gi:6912479 /UG=Hs.45231 leucine zipper, down-regulated in cancer 1 /FL=gb:BC003104.1 gb:AB019527.1 gb:NM\_012317.1

202906\_s\_at Consensus includes gb:AF049895 /DEF=Homo sapiens 8q21.3: Nibrin (NBS1), 2,4-dienoyl-CoA reductase (DECR), and calbindin 1 (CALB1) genes /FEA=mRNA\_5 /DB\_XREF=gi:4126312 /UG=Hs.25812 Nijmegen breakage syndrome 1 (nibrin) /FL=gb:AF058696.1 gb:AF051334.1 gb:NM\_002485.2

**Affy Probe ID**

**Sequence Description**

200743\_s\_at gb:NM\_000391.2 /DEF=Homo sapiens ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) (CLN2), mRNA. /FEA=mRNA /GEN=CLN2 /PROD=ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) /DB\_XREF=gi:5597012 /UG=Hs.20478 ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) /FL=gb:AF017456.1 gb:NM\_000391.2

213620\_s\_at Consensus includes gb:AA126728 /FEA=EST /DB\_XREF=gi:1686292 /DB\_XREF=est:zk95d07.s1 /CLONE=IMAGE:490573 /UG=Hs.83733 intercellular adhesion molecule 2

208729\_x\_at gb:D83043.1 /DEF=Human HLA-B mRNA, allele A\*2711, complete cds. /FEA=mRNA /GEN=HLA-B /DB\_XREF=gi:1871135 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:U36492.1 gb:U35431.1 gb:L37881.1 gb:U31971.1 gb:L42345.1 gb:U50710.1 gb:L41925.1 gb:L41214.1 gb:U27608.1 gb:U35734.1 gb:U70528.1 gb:U70529.1 gb:U80945.1 gb:U64801.1 gb:U09864.1 gb:M77776.1 gb:M77775.1 gb:L04696.1 gb:D83043.1 gb:M12678.1 gb:L09736.1 gb:L09735.1 gb:M75138.1 gb:M84381.1 gb:M84382.1 gb:M84383.1 gb:M84384.1 gb:M84385.1 gb:M32318.1 gb:M32319.1 gb:M32320.1 gb:M81798.1 gb:L04695.1 gb:M84694.1 gb:U30936.1 gb:AF016641.1 gb:U36392.1 gb:AB008102.1 gb:U75533.1 gb:U57966.1 gb:AF026218.1 gb:L07743.1 gb:U01848.1 gb:U38800.1 gb:U40498.1 gb:U03027.1 gb:U03859.1 gb:L22028.1 gb:U04243.1 gb:U06862.1 gb:L32862.1 gb:L15005.1 gb:L22027.1 gb:L19923.1 gb:L11666.1 gb:L11604.1 gb:L11603.1 gb:L19937.1 gb:L33923.1 gb:AF170577.1 gb:M28205.1 gb:U17107.1 gb:U14943.1 gb:AF130734.1 gb:D25275.1 gb:D50290.1 gb:D50291.1 gb:D50299.1 gb:D50300.1 gb:L38504.1 gb:L36591.1 gb:U30904.1

205347\_s\_at gb:NM\_021992.1 /DEF=Homo sapiens thymosin, beta, identified in neuroblastoma cells (TMSNB), mRNA. /FEA=mRNA /GEN=TMSNB /PROD=thymosin, beta, identified in neuroblastoma cells /DB\_XREF=gi:11496272 /UG=Hs.56145 thymosin, beta, identified in neuroblastoma cells /FL=gb:NM\_021992.1 gb:BC000183.1 gb:D82345.1

203504\_s\_at gb:NM\_005502.1 /DEF=Homo sapiens ATP-binding cassette, sub-family A (ABCA1), member 1 (ABCA1), mRNA. /FEA=mRNA /GEN=ABCA1 /PROD=ATP-binding cassette, sub-family A member 1 /DB\_XREF=gi:5915657 /UG=Hs.211562 ATP-binding cassette, sub-family A (ABCA1), member 1 /FL=gb:AF165281.1 gb:NM\_005502.1 gb:AF285167.1

202489\_s\_at gb:BC005238.1 /DEF=Homo sapiens, FXFD domain-containing ion transport regulator 3, clone MGC:12265, mRNA, complete cds. /FEA=mRNA /PROD=FXFD domain-containing ion transport regulator3 /DB\_XREF=gi:13528881 /UG=Hs.301350 FXFD domain-containing ion transport regulator 3 /FL=gb:NM\_005971.2 gb:BC005238.1

221601\_s\_at Consensus includes gb:AI084226 /FEA=EST /DB\_XREF=gi:3422649 /DB\_XREF=est:oy72g09.x1 /CLONE=IMAGE:1671424 /UG=Hs.58831 regulator of Fas-induced apoptosis /FL=gb:AF057557.1 gb:NM\_005449.1

# Affy Probe\_ID

## Sequence Description

221875\_x\_at Consensus includes gb:AW514210 /FEA=EST /DB\_XREF=gi:7152378 /DB\_XREF=est:hd75b05.x1 /CLONE=IMAGE:2915313 /UG=Hs.110309 major histocompatibility complex, class I, F  
219415\_at gb:NM\_020659.1 /DEF=Homo sapiens tweety (Drosophila) homolog 1 (TTYH1), mRNA. /FEA=mRNA /GEN=TTYH1 /PROD=tweety (Drosophila) homolog 1 /DB\_XREF=gi:10257436 /UG=Hs.12828 tweety (Drosophila) homolog 1 /FL=gb:NM\_020659.1 gb:AF177909.1  
207072\_at gb:NM\_003853.1 /DEF=Homo sapiens interleukin 18 receptor accessory protein (IL18RAP), mRNA. /FEA=mRNA /GEN=IL18RAP /PROD=interleukin 18 receptor accessory protein /DB\_XREF=gi:4504656 /UG=Hs.158315 interleukin 18 receptor accessory protein /FL=gb:AF077346.1 gb:NM\_003853.1  
214020\_x\_at Consensus includes gb:BE138575 /FEA=EST /DB\_XREF=gi:8601075 /DB\_XREF=est:xt77e12.x2 /CLONE=IMAGE:2766190 /UG=Hs.149846 integrin, beta 5  
219115\_s\_at gb:NM\_014432.1 /DEF=Homo sapiens interleukin 20 receptor, alpha (IL20RA), mRNA. /FEA=mRNA /GEN=IL20RA /PROD=interleukin 20 receptor, alpha /DB\_XREF=gi:7657690 /UG=Hs.21814 interleukin 20 receptor, alpha /FL=gb:AF184971.1 gb:NM\_014432.1  
204316\_at Consensus includes gb:W19676 /FEA=EST /DB\_XREF=gi:1295644 /DB\_XREF=est:zb36h07.r1 /CLONE=IMAGE:305725 /UG=Hs.82280 regulator of G-protein signalling 10 /FL=gb:NM\_002925.2 gb:AF045229.1  
213075\_at Consensus includes gb:AL050002.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O222 (from clone DKFZp564O222). /FEA=mRNA /DB\_XREF=gi:4884256 /UG=Hs.94795 Homo sapiens mRNA; cDNA DKFZp564O222 (from clone DKFZp564O222)  
213272\_s\_at Consensus includes gb:AF070596.1 /DEF=Homo sapiens clone 24796 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:3387973 /UG=Hs.27191 hypothetical protein from clone 24796  
214368\_at Consensus includes gb:AI68812 /FEA=EST /DB\_XREF=gi:4900106 /DB\_XREF=est:wd41c03.x1 /CLONE=IMAGE:2330692 /UG=Hs.99491 RAS guanyl releasing protein 2 (calcium and DAG-regulated)  
219032\_x\_at gb:NM\_014322.1 /DEF=Homo sapiens opsin 3 (encephalopsin) (OPN3), mRNA. /FEA=mRNA /GEN=OPN3 /PROD=opsin 3 (encephalopsin) /DB\_XREF=gi:7657070 /UG=Hs.279926 opsin 3 (encephalopsin) /FL=gb:AF140242.1 gb:NM\_014322.1  
209344\_at gb:BC002827.1 /DEF=Homo sapiens, tropomyosin 4, clone MGC:3641, mRNA, complete cds. /FEA=mRNA /PROD=tropomyosin 4 /DB\_XREF=gi:12803958 /UG=Hs.250641 tropomyosin 4 /FL=gb:BC002827.1 gb:NM\_003290.1

Affy Probe_ID	Sequence Description
218833_at	gb:NM_016653.1 /DEF=Homo sapiens sterile-alpha motif and leucine zipper containing kinase AZK (ZAK), mRNA. /FEA=mRNA /GEN=ZAK /PROD=sterile-alpha motif and leucine zippercontaining kinase AZK /DB_XREF=gi:7706600 /UG=Hs.115175 sterile-alpha motif and leucine zipper containing kinase AZK /FL=gb:AB049733.1 gb:AF238255.1 gb:AF251441.1 gb:NM_016653.1 gb:AB030034.1
204789_at	gb:NM_005892.1 /DEF=Homo sapiens formin-like (FMNL), mRNA. /FEA=mRNA /GEN=FMNL /PROD=formin-like /DB_XREF=gi:5174400 /UG=Hs.100217 formin-like /FL=gb:BC001710.1 gb:NM_005892.1
218723_s_at	gb:NM_014059.1 /DEF=Homo sapiens RGC32 protein (RGC32), mRNA. /FEA=mRNA /GEN=RGC32 /PROD=RGC32 protein /DB_XREF=gi:7662650 /UG=Hs.76640 RGC32 protein /FL=gb:AF036549.1 gb:NM_014059.1
214071_at	Consensus includes gb:AI082827 /FEA=EST /DB_XREF=gi:3417803 /DB_XREF=est:ox78b09.x1 /CLONE=IMAGE:1662425 /UG=Hs.154145 hypothetical protein FLJ11585
205514_at	gb:NM_018355.1 /DEF=Homo sapiens hypothetical protein FLJ11191 (FLJ11191), mRNA. /FEA=mRNA /GEN=FLJ11191 /PROD=hypothetical protein FLJ11191 /DB_XREF=gi:8922928 /UG=Hs.21838 hypothetical protein FLJ11191 /FL=gb:NM_018355.1
210176_at	gb:AL050262.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I0682 (from clone DKFZp564I0682); complete cds. /FEA=mRNA /GEN=DKFZp564I0682 /PROD=hypothetical protein /DB_XREF=gi:4886482 /UG=Hs.2474 toll-like receptor 1 /FL=gb:U88540.1 gb:D13637.1 gb:NM_003263.1 gb:AL050262.1
213158_at	Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:zk66a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16193 Homo sapiens mRNA; cDNA DKFZp586B211 (from clone DKFZp586B211)
220780_at	gb:NM_015715.1 /DEF=Homo sapiens group III secreted phospholipase A2 (GIII-SPLA2), mRNA. /FEA=mRNA /GEN=GIII-SPLA2 /PROD=group III secreted phospholipase A2 /DB_XREF=gi:7657125 /UG=Hs.149623 group III secreted phospholipase A2 /FL=gb:AF220490.1 gb:NM_015715.1
212185_x_at	Consensus includes gb:NM_005953.1 /DEF=Homo sapiens metallothionein 2A (MT2A), mRNA. /FEA=CDS /GEN=MT2A /PROD=metallothionein 2A /DB_XREF=gi:5174763 /UG=Hs.118786 metallothionein 2A /FL=gb:NM_005953.1
219799_s_at	gb:NM_005771.1 /DEF=Homo sapiens retinol dehydrogenase homolog (RDHL), mRNA. /FEA=mRNA /GEN=RDHL /PROD=retinol dehydrogenase homolog /DB_XREF=gi:5032034 /UG=Hs.179608 retinol dehydrogenase homolog /FL=gb:AF067174.1 gb:NM_005771.1

Affy Probe_ID	Sequence Description
209397_at	gb:BC000147.1 /DEF=Homo sapiens, Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial, clone MGC:5187, mRNA, complete cds. /FEA=mRNA /PROD=Similar to malic enzyme 2, NAD(+)-dependent, mitochondrial /DB_XREF=gi:12652790 /UG=Hs.75342 malic enzyme 2, NAD(+)-dependent, mitochondrial /FL=gb:BC000147.1
218531_at	gb:Nm_025124.1 /DEF=Homo sapiens hypothetical protein FLJ21749 (FLJ21749), mRNA. /FEA=mRNA /GEN=FLJ21749 /PROD=hypothetical protein FLJ21749 /DB_XREF=gi:13376700 /UG=Hs.288761 hypothetical protein FLJ21749 /FL=gb:Nm_025124.1
202747_s_at	gb:Nm_004867.1 /DEF=Homo sapiens integral membrane protein 2A (ITM2A), mRNA. /FEA=mRNA /GEN=ITM2A /PROD=integral membrane protein 2A /DB_XREF=gi:4758223 /UG=Hs.17109 integral membrane protein 2A /FL=gb:AF038953.1 gb:Nm_004867.1
204573_at	gb:Nm_021151.1 /DEF=Homo sapiens carnitine octanoyltransferase (COT), mRNA. /FEA=mRNA /GEN=COT /PROD=carnitine octanoyltransferase /DB_XREF=gi:10863952 /UG=Hs.12743 carnitine O-octanoyltransferase /FL=gb:Nm_021151.1 gb:AF073770.1 gb:AF168793.1
209083_at	gb:U34690.1 /DEF=Human coronin-like protein (HCORO1) mRNA, complete cds. /FEA=mRNA /GEN=HCORO1 /PROD=coronin-like protein /DB_XREF=gi:1002922 /UG=Hs.109606 coronin, actin-binding protein, 1A /FL=gb:U34690.1 gb:Nm_007074.1 gb:D44497.1
220625_s_at	gb:AF115403.1 /DEF=Homo sapiens Ets transcription factor ESE-2b mRNA, complete cds. /FEA=mRNA /PROD=Ets transcription factor ESE-2b /DB_XREF=gi:4559274 /UG=Hs.11713 E74-like factor 5 (ets domain transcription factor) /FL=gb:AF049703.1 gb:Nm_001422.1 gb:AF115403.1
57532_at	Cluster Incl. AW016304:U1-H-BI0p-abl-h-10-0-UI.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-2712427 /clone_end=3 /gb=AW016304 /gi=5865061 /ug=Hs.118640 /len=591
222235_s_at	Consensus includes gb:AL139812 /DEF=Human DNA sequence from clone RP1-19N1 on chromosome Xq21.33-22.3 Contains a gene for a novel protein. Contains ESTs, STSs and GSSs /FEA=mRNA /DB_XREF=gi:9187185 /UG=Hs.287761 Human DNA sequence from clone RP1-19N1 on chromosome Xq21.33-22.3 Contains a gene for a novel protein. Contains ESTs, STSs and GSSs
208121_s_at	gb:Nm_002848.2 /DEF=Homo sapiens protein tyrosine phosphatase, receptor type, O (PTPRO), transcript variant 2, mRNA. /FEA=mRNA /GEN=PTPRO /PROD=receptor-type protein tyrosine phosphatase O, isoform b precursor /DB_XREF=gi:13677212 /FL=gb:Nm_002848.2

# Affy Probe\_ID

## Sequence Description

208986\_at Consensus includes gb:AL559478 /FEA=EST /DB\_XREF=gi:12905019 /DB\_XREF=est:AL559478 /CLONE=CS0DJ013YM17 (5 prime) /UG=Hs.21704 transcription factor 12 (HTF4, helix-loop-helix transcription factors 4) /FL=gb:M80627.1 gb:M83233.1 gb:NM\_003205.2

203490\_at gb:NM\_001421.1 /DEF=Homo sapiens E74-like factor 4 (ets domain transcription factor) (ELF4), mRNA. /FEA=mRNA /GEN=ELF4 /PROD=E74-like factor 4 (ets domain transcriptionfactor) /DB\_XREF=gi:4503554 /UG=Hs.151139 E74-like factor 4 (ets domain transcription factor) /FL=gb:U32645.1 gb:AF000670.1 gb:NM\_001421.1

213119\_at Consensus includes gb:AW058600 /FEA=EST /DB\_XREF=gi:5934239 /DB\_XREF=est:wx23h08.x1 /CLONE=IMAGE:2544543 /UG=Hs.18593 Homo sapiens cDNA: FLJ21449 fis, clone COL04483, highly similar to AF010235 Homo sapiens mRNA from chromosome 5q31-33 region

212314\_at Consensus includes gb:AB018289.1 /DEF=Homo sapiens mRNA for KIAA0746 protein, partial cds. /FEA=mRNA /GEN=KIAA0746 /PROD=KIAA0746 protein /DB\_XREF=gi:3882212 /UG=Hs.49500 KIAA0746 protein

203246\_s\_at gb:NM\_006545.1 /DEF=Homo sapiens homologous to yeast nitrogen permease (candidate tumor suppressor) (NPR2L), mRNA. /FEA=mRNA /GEN=NPR2L /PROD=homologous to yeast nitrogen permease (candidate tumor suppressor) /DB\_XREF=gi:5729948 /UG=Hs.169780 homologous to yeast nitrogen permease (candidate tumor suppressor) /FL=gb:BC002791.1 gb:AF040707.1 gb:NM\_006545.1

205068\_s\_at Consensus includes gb:BE671084 /FEA=EST /DB\_XREF=gi:10031625 /DB\_XREF=est:7e45g09.x1 /CLONE=IMAGE:3285472 /UG=Hs.132942 GTPase regulator associated with the focal adhesion kinase pp125(FAK); KIAA0621 protein /FL=gb:AF141884.1 gb:NM\_015071.1

210790\_s\_at gb:BC003658.1 /DEF=Homo sapiens, SAR1 protein, clone MGC:4824, mRNA, complete cds. /FEA=mRNA /PROD=SAR1 protein /DB\_XREF=gi:1317777 /UG=Hs.110796 SAR1 protein /FL=gb:BC003658.1

204793\_at gb:NM\_014710.1 /DEF=Homo sapiens KIAA0443 gene product (KIAA0443), mRNA. /FEA=mRNA /GEN=KIAA0443 /PROD=KIAA0443 gene product /DB\_XREF=gi:7662129 /UG=Hs.113082 KIAA0443 gene product /FL=gb:AB007903.1 gb:NM\_014710.1

205475\_at gb:NM\_007281.1 /DEF=Homo sapiens scrapie responsive protein 1 (SCRG1), mRNA. /FEA=mRNA /GEN=SCRG1 /PROD=scrapie responsive protein 1 /DB\_XREF=gi:6005869 /UG=Hs.7122 scrapie responsive protein 1 /FL=gb:NM\_007281.1

203074\_at gb:NM\_001630.1 /DEF=Homo sapiens annexin A8 (ANXA8), mRNA. /FEA=mRNA /GEN=ANXA8 /PROD=annexin VIII /DB\_XREF=gi:4502112 /UG=Hs.87268 annexin A8 /FL=gb:BC004376.1 gb:M81844.1 gb:NM\_001630.1

# Affy Probe\_ID

## Sequence Description

204140\_at gb:NM\_003596.1 /DEF=Homo sapiens tyrosylprotein sulfotransferase 1 (TPST1), mRNA. /FEA=mRNA /GEN=TPST1 /PROD=tyrosylprotein sulfotransferase 1 /DB\_XREF=gi:4507664 /UG=Hs.17279 tyrosylprotein sulfotransferase 1 /FL=gb:AF038009.1 gb:NM\_003596.1

214435\_x\_at Consensus includes gb:NM\_005402.1 /DEF=Homo sapiens v-ral simian leukemia viral oncogene homolog A (ras related) (RALA), mRNA. /FEA=CDS /GEN=RALA /PROD=v-ral simian leukemia viral oncogene homolog A (ras related) /DB\_XREF=gi:4885568 /UG=Hs.288757 v-ral simian leukemia viral oncogene homolog A (ras related) /FL=gb:M29893.1 gb:NM\_005402.1

200967\_at gb:NM\_000942.1 /DEF=Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB), mRNA. /FEA=mRNA /GEN=PPIB /PROD=peptidylprolyl isomerase B (cyclophilin B) /DB\_XREF=gi:4758949 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin B) /FL=gb:BC001125.1 gb:M60857.1 gb:NM\_000942.1

202304\_at gb:NM\_014923.1 /DEF=Homo sapiens KIAA0970 protein (KIAA0970), mRNA. /FEA=mRNA /GEN=KIAA0970 /PROD=KIAA0970 protein /DB\_XREF=gi:7662419 /UG=Hs.103329 KIAA0970 protein /FL=gb:AB023187.1 gb:NM\_014923.1

216742\_at Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fis, clone ADSE00417. /FEA=mRNA /DB\_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens cDNA: FLJ20908 fis, clone ADSE00417

204150\_at gb:NM\_015136.1 /DEF=Homo sapiens KIAA0246 protein (KIAA0246), mRNA. /FEA=mRNA /GEN=KIAA0246 /PROD=KIAA0246 protein /DB\_XREF=gi:12225239 /UG=Hs.301989 KIAA0246 protein /FL=gb:NM\_015136.1

204897\_at Consensus includes gb:AA897516 /FEA=EST /DB\_XREF=gi:3034136 /DB\_XREF=est:aj62c04.s1 /CLONE=IMAGE:1394886 /UG=Hs.199248 prostaglandin E receptor 4 (subtype EP4) /FL=gb:D28472.1 gb:L25124.1 gb:NM\_000958.1 gb:L28175.1

202669\_s\_at gb:U16797.1 /DEF=Human LERK-5 (EPLG5) mRNA, complete cds. /FEA=mRNA /GEN=EPLG5 /PROD=LERK-5 /DB\_XREF=gi:902370 /UG=Hs.30942 ephrin-B2 /FL=gb:U81262.1 gb:NM\_004093.1 gb:L38734.1 gb:U16797.1

212909\_at Consensus includes gb:AL567376 /FEA=EST /DB\_XREF=gi:12920672 /DB\_XREF=est:AL567376 /CLONE=CS0DF024YO04 (3 prime) /UG=Hs.85339 G protein-coupled receptor 39

211163\_s\_at gb:AF012536.1 /DEF=Homo sapiens decoy receptor 1 (DcR1) mRNA, complete cds. /FEA=mRNA /GEN=DcR1 /PROD=decoy receptor 1 /DB\_XREF=gi:2338421 /UG=Hs.119684 tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain /FL=gb:AF012536.1

Affy Probe_ID	Sequence Description
212531_at	Consensus includes gb:NM_005564.1 /DEF=Homo sapiens lipocalin 2 (oncogene 24p3) (LCN2), mRNA. /FEA=CDS /GEN=LCN2 /PROD=lipocalin 2 (oncogene 24p3) /DB_XREF=gi:5031852 /UG=Hs.204238 lipocalin 2 (oncogene 24p3) /FL=gb:NM_005564.1
205639_at	gb:NM_001637.1 /DEF=Homo sapiens acylxyacyl hydrolase (neutrophil) (AOAH), mRNA. /FEA=mRNA /GEN=AOAH /PROD=acylxyacyl hydrolase precursor /DB_XREF=gi:4502114 /UG=Hs.82542 acylxyacyl hydrolase (neutrophil) /FL=gb:M62840.1 gb:NM_001637.1
214175_x_at	Consensus includes gb:AI254547 /FEA=EST /DB_XREF=gi:3862072 /DB_XREF=est:qv48h06.x1 /CLONE=IMAGE:1984859 /UG=Hs.79691 LIM domain protein
201674_s_at	gb:BC000729.1 /DEF=Homo sapiens, A kinase (PRKA) anchor protein 1, clone MGC:1807, mRNA, complete cds. /FEA=mRNA /PROD=A kinase (PRKA) anchor protein 1 /DB_XREF=gi:12653874 /UG=Hs.78921 A kinase (PRKA) anchor protein 1 /FL=gb:BC000729.1 gb:NM_003488.1
205795_at	gb:NM_004796.1 /DEF=Homo sapiens neurexin 3 (NRXN3), mRNA. /FEA=mRNA /GEN=NRXN3 /PROD=neurexin 3 /DB_XREF=gi:4758827 /UG=Hs.22269 neurexin 3 /FL=gb:AB018286.1 gb:NM_004796.1
203066_at	gb:NM_014863.1 /DEF=Homo sapiens B cell RAG associated protein (BRAG), mRNA. /FEA=mRNA /GEN=BRAG /PROD=KIAA0598 gene product /DB_XREF=gi:7662195 /UG=Hs.6079 B cell RAG associated protein /FL=gb:AB011170.1 gb:AF026477.1 gb:NM_014863.1 gb:NM_015892.1
219401_at	gb:NM_022167.1 /DEF=Homo sapiens xylosyltransferase II (XT2), mRNA. /FEA=mRNA /GEN=XT2 /PROD=xylosyltransferase II /DB_XREF=gi:11545913 /UG=Hs.32117 xylosyltransferase II /FL=gb:NM_022167.1
205401_at	gb:NM_003659.1 /DEF=Homo sapiens alkylglycerone phosphate synthase (AGPS), mRNA. /FEA=mRNA /GEN=AGPS /PROD=alkylglycerone phosphate synthase precursor /DB_XREF=gi:4501992 /UG=Hs.22580 alkylglycerone phosphate synthase /FL=gb:NM_003659.1
202756_s_at	gb:NM_002081.1 /DEF=Homo sapiens glypican 1 (GPC1), mRNA. /FEA=mRNA /GEN=GPC1 /PROD=glypican 1 precursor /DB_XREF=gi:4504080 /UG=Hs.2699 glypican 1 /FL=gb:NM_002081.1
213874_at	Consensus includes gb:NM_006215.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 4 (SERPINA4), mRNA. /FEA=CDS /GEN=SERPINA4 /PROD=serine (or cysteine) proteinase inhibitor, cladeA (alpha-1 antiproteinase, antitrypsin), member 4 /DB_XREF=gi:5453887 /UG=Hs.159628 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 4 /FL=gb:L19684.1 gb:NM_006215.1



Affy Probe_ID	Sequence Description
219100_at	gb:NM_024928.1 /DEF=Homo sapiens hypothetical protein FLJ22559 (FLJ22559), mRNA. /FEA=mRNA /GEN=FLJ22559 /PROD=hypothetical protein FLJ22559 /DB_XREF=gi:13487922 /UG=Hs.273387 hypothetical protein FLJ22559 /FL=gb:NM_024928.1
205388_at	gb:NM_003279.1 /DEF=Homo sapiens troponin C2, fast (TNNC2), mRNA. /FEA=mRNA /GEN=TNNC2 /PROD=troponin C2, fast /DB_XREF=gi:4507616 /UG=Hs.182421 troponin C2, fast /FL=gb:M22307.1 gb:BC005323.1 gb:NM_003279.1
206039_at	gb:NM_004794.1 /DEF=Homo sapiens RAB33A, member RAS oncogene family (RAB33A), mRNA. /FEA=mRNA /GEN=RAB33A /PROD=RAB33A, member RAS oncogene family /DB_XREF=gi:4758995 /UG=Hs.56294 RAB33A, member RAS oncogene family /FL=gb:BC001157.1 gb:D14889.1 gb:NM_004794.1 Consensus includes gb:AI927770 /FEA=EST /DB_XREF=gi:5663734 /DB_XREF=est:wo08h07.x1 /CLONE=IMAGE:2454781 /UG=Hs.181300 sel-1 (suppressor of lin-12, C.elegans)-like /FL=gb:NM_005065.1 gb:AB020335.1 gb:AF052059.1 gb:U11037.1
202061_s_at	gb:NM_006518.1 /DEF=Homo sapiens small proline-rich protein 2C (SPRR2C), mRNA. /FEA=mRNA /GEN=SPRR2C /PROD=small proline-rich protein 2C /DB_XREF=gi:10835253 /UG=Hs.2421 small proline-rich protein 2C /FL=gb:NM_006518.1
220664_at	gb:NM_025176.1 /DEF=Homo sapiens KIAA0980 protein (KIAA0980), mRNA. /FEA=mRNA /GEN=KIAA0980 /PROD=hypothetical protein FLJ11792 /DB_XREF=gi:13378150 /UG=Hs.227743 KIAA0980 protein /FL=gb:NM_025176.1
212829_at	Consensus includes gb:BE878277 /FEA=EST /DB_XREF=gi:10327053 /DB_XREF=est:601488188F1 /CLONE=IMAGE:3890564 /UG=Hs.57079 Homo sapiens cDNA FLJ13267 fis, clone OVARC1000964 gb:L42024.1 /DEF=Homo sapiens MHC HLA-B39 mRNA, complete cds. /FEA=mRNA /GEN=HLA-B39 /PROD=major histocompatibility complex /DB_XREF=gi:804748 /UG=Hs.77961 major histocompatibility complex, class I, B /FL=gb:L37880.1 gb:U29057.1 gb:U29480.1 gb:U63653.1 gb:D85761.1 gb:D85762.1 gb:M77774.1 gb:M77778.1 gb:M77777.1 gb:M16102.1 gb:M84380.1 gb:M32317.1 gb:M24033.1 gb:L17005.1 gb:U88407.1 gb:U29083.1 gb:L24373.1 gb:U03698.1 gb:U04787.1 gb:U04244.1 gb:U04245.1 gb:NM_005514.1 gb:L33922.1 gb:AF189017.1 gb:U21052.1 gb:U21053.1 gb:L36318.1 gb:U09912.1 gb:L42024.1
203595_s_at	Consensus includes gb:N47725 /FEA=EST /DB_XREF=gi:1188891 /DB_XREF=est:yy92h11.s1 /CLONE=IMAGE:281061 /UG=Hs.27610 retinoic acid- and interferon-inducible protein (58kD) /FL=gb:U34605.1 gb:NM_012420.1

# Affy Probe\_ID

## Sequence Description

204780_s_at	Consensus includes gb:AA164751 /FEA=EST /DB_XREF=gi:1740929 /DB_XREF=est:zo93g12.s1 /CLONE=IMAGE:594502 /UG=Hs.82359 tumor necrosis factor receptor superfamily, member 6 /FL=gb:M67454.1 gb:NM_0000043.1
202481_at	gb:NM_004753.1 /DEF=Homo sapiens short-chain dehydrogenasereductase 1 (SDR1), mRNA. /FEA=mRNA /GEN=SDR1 /PROD=short-chain dehydrogenasereductase 1 /DB_XREF=gi:4759083 /UG=Hs.17144 short-chain dehydrogenasereductase 1 /FL=gb:BC002730.1 gb:AF061741.1 gb:NM_004753.1
208072_s_at	gb:NM_003648.1 /DEF=Homo sapiens diacylglycerol kinase, delta (130kD) (DGKD), mRNA. /FEA=mRNA /GEN=DGKD /PROD=diacylglycerol kinase, delta (130kD) /DB_XREF=gi:4503310 /UG=Hs.115907 diacylglycerol kinase, delta (130kD) /FL=gb:D73409.1 gb:NM_003648.1
217502_at	Consensus includes gb:BE888744 /FEA=EST /DB_XREF=gi:10345354 /DB_XREF=est:601513656F1 /CLONE=IMAGE:3915133 /UG=Hs.169274 ESTs, Highly similar to IFT2_HUMAN INTERFERON-INDUCED PROTEIN WITH TETRATRICHOPEPTIDE REPEATS 2 H.sapiens
205637_s_at	gb:NM_003027.1 /DEF=Homo sapiens SH3-domain GRB2-like 3 (SH3GL3), mRNA. /FEA=mRNA /GEN=SH3GL3 /PROD=SH3-domain GRB2-like 3 /DB_XREF=gi:4506932 /UG=Hs.80315 SH3-domain GRB2-like 3 /FL=gb:AF036269.1 gb:AF036270.1 gb:AF036271.1 gb:NM_003027.1
219732_at	gb:NM_017753.1 /DEF=Homo sapiens hypothetical protein FLJ20300 (FLJ20300), mRNA. /FEA=mRNA /GEN=FLJ20300 /PROD=hypothetical protein FLJ20300 /DB_XREF=gi:8923280 /UG=Hs.106825 hypothetical protein FLJ20300 /FL=gb:NM_017753.1
202877_s_at	Consensus includes gb:W72082 /FEA=EST /DB_XREF=gi:1382588 /DB_XREF=est:zd70c06.s1 /CLONE=IMAGE:345994 /UG=Hs.97199 complement component C1q receptor /FL=gb:NM_012072.2 gb:U94333.1
215790_at	Consensus includes gb:AA835004 /FEA=EST /DB_XREF=gi:2908732 /DB_XREF=est:aj39d06.s1 /CLONE=IMAGE:1392683 /UG=Hs.25924 hypothetical protein
214459_x_at	Consensus includes gb:M12679.1 /DEF=Human Cw1 antigen mRNA, complete cds. /FEA=mRNA /GEN=HLA-C /DB_XREF=gi:187911 /UG=Hs.274485 Cw1 antigen /FL=gb:M12679.1
202458_at	gb:NM_007173.1 /DEF=Homo sapiens protease, serine, 23 (SPUVE), mRNA. /FEA=mRNA /GEN=SPUVE /PROD=protease, serine, 23 /DB_XREF=gi:6005881 /UG=Hs.325820 protease, serine, 23 /FL=gb:AL136914.1 gb:BC001278.1 gb:AF015287.1 gb:NM_007173.1 gb:AF193611.1

Affy Probe_ID	Sequence Description
204675_at	gb:NM_001047.1 /DEF=Homo sapiens steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1) (SRD5A1), mRNA. /FEA=mRNA /GEN=SRD5A1 /PROD=steroid-5-alpha-reductase, alpha polypeptide 1(3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1) /DB_XREF=gi:4507200 /UG=Hs.552 steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1) /FL=gb:M32313.1 gb:AF052126.1 gb:NM_001047.1
203764_at	gb:NM_014750.1 /DEF=Homo sapiens KIAA0008 gene product (KIAA0008), mRNA. /FEA=mRNA /GEN=KIAA0008 /PROD=KIAA0008 gene product /DB_XREF=gi:7661851 /UG=Hs.77695 KIAA0008 gene product /FL=gb:D13633.1 gb:NM_014750.1
202557_at	Consensus includes gb:AI718418 /FEA=EST /DB_XREF=gi:5035674 /DB_XREF=est:as43b01.x1 /CLONE=IMAGE:2319913 /UG=Hs.288799 stress 70 protein chaperone, microsome-associated, 60kD /FL=gb:U04735.1 gb:NM_006948.1
220104_at	gb:NM_020119.1 /DEF=Homo sapiens hypothetical protein FLB6421 (FLB6421), mRNA. /FEA=mRNA /GEN=FLB6421 /PROD=hypothetical protein FLB6421 /DB_XREF=gi:9910221 /UG=Hs.35254 hypothetical protein FLB6421 /FL=gb:NM_020119.1
205259_at	gb:NM_000901.1 /DEF=Homo sapiens nuclear receptor subfamily 3, group C, member 2 (NR3C2), mRNA. /FEA=mRNA /GEN=NR3C2 /PROD=nuclear receptor subfamily 3, group C, member 2 /DB_XREF=gi:4505198 /UG=Hs.1790 nuclear receptor subfamily 3, group C, member 2 /FL=gb:M16801.1 gb:NM_000901.1
206167_s_at	gb:NM_001174.2 /DEF=Homo sapiens Rho GTPase activating protein 6 (ARHGAP6), transcript variant 2, mRNA. /FEA=mRNA /GEN=ARHGAP6 /PROD=Rho GTPase activating protein 6 isoform 2 /DB_XREF=gi:7382476 /UG=Hs.250830 Rho GTPase activating protein 6 /FL=gb:AF022212.2 gb:NM_001174.2
217567_at	Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:11129195 /DB_XREF=est:7p41a04.x1 /CLONE=IMAGE:3648271 /UG=Hs.289803 ESTs, Moderately similar to S71105 protein-glutamine gamma-glutamyltransferase H.sapiens
208712_at	gb:M73554.1 /DEF=Human bcl-1 mRNA, complete CDS. /FEA=mRNA /GEN=bcl-1 /PROD=bcl-1 /DB_XREF=gi:179364 /UG=Hs.82932 cyclin D1 (PRAD1: parathyroid adenomatosis 1) /FL=gb:BC000076.1 gb:M73554.1
219429_at	gb:NM_024306.1 /DEF=Homo sapiens fatty acid hydroxylase (FAAH), mRNA. /FEA=mRNA /GEN=FAAH /PROD=fatty acid hydroxylase /DB_XREF=gi:13236537 /UG=Hs.249163 fatty acid hydroxylase /FL=gb:BC002679.1 gb:NM_024306.1 gb:BC0004263.1

# Affy Probe ID

## Sequence Description

205844\_at gb:NM\_004666.1 /DEF=Homo sapiens vanin 1 (VNN1), mRNA. /FEA=mRNA /GEN=VNN1 /PROD=vanin 1 /DB\_XREF=gi:4759311 /UG=Hs.12114 vanin 1 /FL=gb:NM\_004666.1 gb:U39664.1  
203642\_s\_at gb:NM\_014900.1 /DEF=Homo sapiens KIAA0977 protein (KIAA0977), mRNA. /FEA=mRNA /GEN=KIAA0977 /PROD=KIAA0977 protein /DB\_XREF=gi:7662427 /UG=Hs.300855 KIAA0977 protein /FL=gb:AB023194.1 gb:NM\_014900.1  
205126\_at gb:NM\_006296.1 /DEF=Homo sapiens vaccinia related kinase 2 (VRK2), mRNA. /FEA=mRNA /GEN=VRK2 /PROD=vaccinia related kinase 2 /DB\_XREF=gi:5454163 /UG=Hs.82771 vaccinia related kinase 2 /FL=gb:AB000450.1 gb:NM\_006296.1  
203921\_at gb:NM\_004267.1 /DEF=Homo sapiens carbohydrate (chondroitin 6keratan) sulfotransferase 2 (CHST2), mRNA. /FEA=mRNA /GEN=CHST2 /PROD=carbohydrate (chondroitin 6keratan)sulfotransferase 2 /DB\_XREF=gi:4757983 /UG=Hs.8786 carbohydrate (chondroitin 6keratan) sulfotransferase 2 /FL=gb:AB021124.1 gb:AB014679.1 gb:AB014680.1 gb:AF083066.1 gb:NM\_004267.1  
213274\_s\_at Consensus includes gb:AA020826 /FEA=EST /DB\_XREF=gi:1484570 /DB\_XREF=est:ze64b04.sl /CLONE=IMAGE:363727 /UG=Hs.297939 cathepsin B  
221058\_s\_at gb:NM\_016326.2 /DEF=Homo sapiens chemokine-like factor 3, alternatively spliced (LOC51192), mRNA. /FEA=mRNA /GEN=LOC51192 /PROD=chemokine-like factor 3, alternatively spliced /DB\_XREF=gi:10092611 /UG=Hs.15159 chemokine-like factor, alternatively spliced /FL=gb:NM\_016326.2 gb:AF135381.2  
214451\_at Consensus includes gb:NM\_003221.1 /DEF=Homo sapiens transcription factor AP-2 beta (activating enhancer-binding protein 2 beta) (TFAP2B), mRNA. /FEA=CDS /GEN=TFAP2B /PROD=transcription factor AP-2 beta (activating enhancer-binding protein 2 beta) /DB\_XREF=gi:4507442 /UG=Hs.33102 transcription factor AP-2 beta (activating enhancer-binding protein 2 beta) /FL=gb:NM\_003221.1  
207254\_at gb:NM\_005073.1 /DEF=Homo sapiens solute carrier family 15 (oligopeptide transporter), member 1 (SLC15A1), mRNA. /FEA=mRNA /GEN=SLC15A1 /PROD=solute carrier family 15 (oligopeptidetransporter), member 1 /DB\_XREF=gi:4827007 /UG=Hs.2217 solute carrier family 15 (oligopeptide transporter), member 1 /FL=gb:AF043233.1 gb:NM\_005073.1 gb:U21936.1 gb:U13173.1  
207565\_s\_at gb:NM\_001531.1 /DEF=Homo sapiens major histocompatibility complex, class I-like sequence (HLALS), mRNA. /FEA=mRNA /GEN=HLALS /PROD=major histocompatibility complex, class I-like sequence /DB\_XREF=gi:4504416 /UG=Hs.101840 major histocompatibility complex, class I-like sequence /FL=gb:NM\_001531.1 gb:U22963.1

**Affy Probe ID**

**Sequence Description**

213939_s_at	Consensus includes gb:AI871641 /FEA=EST /DB_XREF=gi:5545690 /DB_XREF=est:we28f04.x1 /CLONE=IMAGE:2342431 /UG=Hs.306623 Homo sapiens cDNA FLJ12012 fs, clone HEMBB1001668
208998_at	gb:U94592.1 /DEF=Human uncoupling protein homolog (UCPH) mRNA, complete cds. /FEA=mRNA /GEN=UCPH /PROD=uncoupling protein homolog /DB_XREF=gi:2052354 /UG=Hs.80658 uncoupling protein 2 (mitochondrial, proton carrier) /FL=gb:NM_003355.2 gb:U76367.1 gb:U82819.1 gb:U94592.1
221665_s_at	gb:BC004907.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 4632407K17 gene, clone MGC:4642, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 4632407K17 gene /DB_XREF=gi:13436193 /UG=Hs.28907 hypothetical protein FLJ20258 /FL=gb:BC004907.1
216231_s_at	Consensus includes gb:AW188940 /FEA=EST /DB_XREF=gi:6463376 /DB_XREF=est:xl90f09.x1 /CLONE=IMAGE:2674025 /UG=Hs.75415 beta-2-microglobulin
200958_s_at	gb:NM_005625.1 /DEF=Homo sapiens syndecan binding protein (syntenin) (SDCBP), mRNA. /FEA=mRNA /GEN=SDCBP /PROD=syndecan binding protein (syntenin) /DB_XREF=gi:5032082 /UG=Hs.8180 syndecan binding protein (syntenin) /FL=gb:AF000652.1 gb:NM_005625.1
202644_s_at	gb:NM_006290.1 /DEF=Homo sapiens tumor necrosis factor, alpha-induced protein 3 (TNFAIP3), mRNA. /FEA=mRNA /GEN=TNFAIP3 /PROD=tumor necrosis factor, alpha-induced protein 3 /DB_XREF=gi:5454131 /UG=Hs.211600 tumor necrosis factor, alpha-induced protein 3 /FL=gb:M59465.1 gb:NM_006290.1
206995_x_at	gb:NM_003693.1 /DEF=Homo sapiens acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells (SREC), mRNA. /FEA=mRNA /GEN=SREC /PROD=acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells /DB_XREF=gi:4507202 /UG=Hs.57735 acetyl LDL receptor; SREC /FL=gb:D63483.1 gb:D86864.1 gb:NM_003693.1
218935_at	gb:NM_014600.1 /DEF=Homo sapiens EH-domain containing 3 (EHD3), mRNA. /FEA=mRNA /GEN=EHD3 /PROD=EH-domain containing 3 /DB_XREF=gi:7657055 /UG=Hs.87125 EH-domain containing 3 /FL=gb:AF214736.1 gb:AF181264.1 gb:NM_014600.1
210314_x_at	gb:AF114013.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1 gamma mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related deathligand-1 gamma /DB_XREF=gi:7328557 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF136294.1 gb:AF114013.1
219334_s_at	gb:NM_022837.1 /DEF=Homo sapiens hypothetical protein FLJ22833 (FLJ22833), mRNA. /FEA=mRNA /GEN=FLJ22833 /PROD=hypothetical protein FLJ22833 /DB_XREF=gi:12383083 /UG=Hs.118183 hypothetical protein FLJ22833 /FL=gb:NM_022837.1

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## Sequence Description

205005\_s\_at Consensus includes gb:AW293531 /FEA=EST /DB\_XREF=gi:6700167 /DB\_XREF=est:UI-H-BI2-ahq-d-03-0-UI.s1 /CLONE=IMAGE:2727604 /UG=Hs.122647 N-myristoyltransferase 2 /FL=gb:AF043325.1 gb:NM\_004808.1 gb:AF280094.1 /DEF=Homo sapiens transcriptional coactivator Sp110b mRNA, complete cds. /FEA=mRNA /PROD=transcriptional coactivator Sp110b /DB\_XREF=gi:9800493 /UG=Hs.38125 interferon-induced protein 75, 52kD /FL=gb:AF280094.1

217648\_at Consensus includes gb:AW295367 /FEA=EST /DB\_XREF=gi:6702003 /DB\_XREF=est:UI-H-BI2-ahv-e-04-0-UI.s1 /CLONE=IMAGE:2728039 /UG=Hs.196585 ESTs, Weakly similar to Pro-Pol-dUTPase polyprotein M.musculus

201329\_s\_at gb:NM\_005239.1 /DEF=Homo sapiens v-ets avian erythroblastosis virus E26 oncogene homolog 2 (ETS2), mRNA. /FEA=mRNA /GEN=ETS2 /PROD=v-ets avian erythroblastosis virus E26 oncogene homolog 2 /DB\_XREF=gi:4885220 /UG=Hs.85146 v-ets avian erythroblastosis virus E26 oncogene homolog 2 /FL=gb:J04102.1 gb:NM\_005239.1

202270\_at gb:NM\_002053.1 /DEF=Homo sapiens guanylate binding protein 1, interferon-inducible, 67kD (GBP1), mRNA. /FEA=mRNA /GEN=GBP1 /PROD=guanylate binding protein 1,interferon-inducible, 67kD /DB\_XREF=gi:4503938 /UG=Hs.62661 guanylate binding protein 1, interferon-inducible, 67kD /FL=gb:BC002666.1 gb:M55542.1 gb:NM\_002053.1

215908\_at Consensus includes gb:AF009267.1 /DEF=Homo sapiens clone FBA1 Cri-du-chat region mRNA. /FEA=mRNA /DB\_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens clone FBA1 Cri-du-chat region mRNA

215146\_s\_at Consensus includes gb:AB028966.1 /DEF=Homo sapiens mRNA for KIAA1043 protein, partial cds. /FEA=mRNA /GEN=KIAA1043 /PROD=KIAA1043 protein /DB\_XREF=gi:5689422 /UG=Hs.11390 KIAA1043 protein

218259\_at gb:NM\_014048.1 /DEF=Homo sapiens KIAA1243 protein (KIAA1243), mRNA. /FEA=mRNA /GEN=KIAA1243 /PROD=NPD001 protein /DB\_XREF=gi:7662513 /UG=Hs.151076 KIAA1243 protein /FL=gb:AF078853.1 gb:NM\_014048.1

209759\_s\_at gb:BC002746.1 /DEF=Homo sapiens, Similar to dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase), clone MGC:3903, mRNA, complete cds. /FEA=mRNA /PROD=Similar to dodecenoyl-Coenzyme A delta isomerase(3,2 trans-enoyl-Coenzyme A isomerase) /DB\_XREF=gi:12803812 /UG=Hs.322824 Homo sapiens, Similar to dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase), clone MGC:3903, mRNA, complete cds /FL=gb:BC002746.1

# Affy Probe\_ID

## Sequence Description

218440\_at gb:NM\_020166.2 /DEF=Homo sapiens methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) (MCCC1), mRNA. /FEA=mRNA /GEN=MCCC1 /PROD=methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) /DB\_XREF=gi:13518227 /UG=Hs.47649 methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) /FL=gb:AF310339.1 gb:AF310972.1

209392\_at gb:BC004187.1 gb:BC004214.1 gb:NM\_020166.2 gb:AB029826.1

202803\_s\_at gb:L35594.1 /DEF=Human autotaxin mRNA, complete cds. /FEA=mRNA /PROD=autotaxin /DB\_XREF=gi:537905 /UG=Hs.174185 ectonucleotide pyrophosphatase phosphodiesterase 2 (autotaxin) /FL=gb:L35594.1

202803\_s\_at gb:NM\_000211.1 /DEF=Homo sapiens integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) (ITGB2), mRNA. /FEA=mRNA /GEN=ITGB2 /PROD=integrin beta chain, beta 2 precursor /DB\_XREF=gi:4557885 /UG=Hs.83968 integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) /FL=gb:NM\_000211.1

219377\_at gb:NM\_022751.1 /DEF=Homo sapiens hypothetical protein FLJ21610 (FLJ21610), mRNA. /FEA=mRNA /GEN=FLJ21610 /PROD=hypothetical protein FLJ21610 /DB\_XREF=gi:12232414 /UG=Hs.12727 hypothetical protein FLJ21610 /FL=gb:NM\_022751.1

218671\_s\_at gb:NM\_016311.1 /DEF=Homo sapiens ATPase inhibitor precursor (LOC511189), mRNA. /FEA=mRNA /GEN=LOC511189 /PROD=ATPase inhibitor precursor /DB\_XREF=gi:7705926 /UG=Hs.241336 ATPase inhibitor precursor /FL=gb:BC004955.1 gb:AB029042.1 gb:NM\_016311.1

201337\_s\_at gb:NM\_004781.2 /DEF=Homo sapiens vesicle-associated membrane protein 3 (cellubrevin) (VAMP3), mRNA. /FEA=mRNA /GEN=VAMP3 /PROD=vesicle-associated membrane protein 3 /DB\_XREF=gi:9257252 /UG=Hs.66708 vesicle-associated membrane protein 3 (cellubrevin) /FL=gb:BC003570.1 gb:NM\_004781.2

204786\_s\_at gb:L41944.1 /DEF=Homo sapiens interferon receptor ifnar2-1 (splice variant IFNAR2-1) mRNA, complete cds. /FEA=mRNA /GEN=IFNAR2 /PROD=interferon receptor /DB\_XREF=gi:995296 /UG=Hs.86958 interferon (alpha, beta and omega) receptor 2 /FL=gb:NM\_000874.1 gb:L41944.1

203021\_at gb:NM\_003064.1 /DEF=Homo sapiens secretory leukocyte protease inhibitor (antileukoproteinase) (SLPI), mRNA. /FEA=mRNA /GEN=SLPI /PROD=secretory leukocyte protease inhibitor (antileukoproteinase) /DB\_XREF=gi:4507064 /UG=Hs.251754 secretory leukocyte protease inhibitor (antileukoproteinase) /FL=gb:NM\_003066.1 gb:AF114471.1 gb:NM\_003064.1

214211\_at Consensus includes gb:AA083483 /FEA=EST /DB\_XREF=gi:1625544 /DB\_XREF=est:zn32a06.s1. /CLONE=IMAGE:549106 /UG=Hs.62954 ferritin, heavy polypeptide 1

# Affy Probe ID

## Sequence Description

218876\_at gb:NM\_016140.1 /DEF=Homo sapiens brain specific protein (LOC51673), mRNA. /FEA=mRNA /GEN=LOC51673 /PROD=brain specific protein /DB\_XREF=gi:7706392 /UG=Hs.279772 brain specific protein /FL=gb:BC000691.1 gb:AF132972.1 gb:AF078846.1 gb:NM\_015964.1 gb:NM\_016140.1

201089\_at gb:NM\_001693.1 /DEF=Homo sapiens ATPase, H<sup>+</sup> transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 (ATP6B2), mRNA. /FEA=mRNA /GEN=ATP6B2 /PROD=ATPase, H<sup>+</sup> transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 /DB\_XREF=gi:4502310 /UG=Hs.1697 ATPase, H<sup>+</sup> transporting, lysosomal (vacuolar proton pump), beta polypeptide, 5658kD, isoform 2 /FL=gb:BC003100.1 gb:NM\_001693.1 gb:L35249.1

213638\_at Consensus includes gb:AW054711 /FEA=EST /DB\_XREF=gi:5920414 /DB\_XREF=est:wz96g08.x1 /CLONE=IMAGE:2566718 /UG=Hs.121591 Human DNA sequence from PAC 257A7 on chromosome 6p24. Contains two unknown genes and ESTs, STSs and a GSS

206130\_s\_at gb:NM\_001181.1 /DEF=Homo sapiens asialoglycoprotein receptor 2 (ASGR2), mRNA. /FEA=mRNA /GEN=ASGR2 /PROD=asialoglycoprotein receptor 2 /DB\_XREF=gi:4502252 /UG=Hs.1259 asialoglycoprotein receptor 2 /FL=gb:M11025.1 gb:NM\_001181.1

205014\_at gb:NM\_005130.1 /DEF=Homo sapiens heparin-binding growth factor binding protein (HBP17), mRNA. /FEA=mRNA /GEN=HBP17 /PROD=heparin-binding growth factor binding protein /DB\_XREF=gi:4826753 /UG=Hs.1690 heparin-binding growth factor binding protein /FL=gb:BC003628.1 gb:M60047.1 gb:NM\_005130.1

221713\_s\_at gb:BC006434.1 /DEF=Homo sapiens, hypothetical protein FLJ12748, clone MGC:13089, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ12748 /DB\_XREF=gi:13623628 /FL=gb:BC006434.1

206683\_at gb:NM\_003447.1 /DEF=Homo sapiens zinc finger protein 165 (ZNF165), mRNA. /FEA=mRNA /GEN=ZNF165 /PROD=zinc finger protein 165 /DB\_XREF=gi:4508000 /UG=Hs.55481 zinc finger protein 165 /FL=gb:U78722.1 gb:NM\_003447.1

202084\_s\_at gb:NM\_003003.1 /DEF=Homo sapiens SEC14 (S. cerevisiae)-like 1 (SEC14L1), mRNA. /FEA=mRNA /GEN=SEC14L1 /PROD=SEC14 (S. cerevisiae)-like 1 /DB\_XREF=gi:4506866 /UG=Hs.75232 SEC14 (S. cerevisiae)-like 1 /FL=gb:D67029.1 gb:NM\_003003.1

211694\_at gb:AF348076.1 /DEF=Homo sapiens serinethreonine kinase FKSG81 (FKSG81) mRNA, complete cds. /FEA=mRNA /GEN=FKSG81 /PROD=serinethreonine kinase FKSG81 /DB\_XREF=gi:13540323 /FL=gb:AF348076.1

212481\_s\_at Consensus includes gb:AI214061 /FEA=EST /DB\_XREF=gi:3777662 /DB\_XREF=est:ap29d10.x1 /CLONE=IMAGE:1956787 /UG=Hs.250641 tropomyosin 4



Affy Probe ID	Sequence Description
205796_at	gb:NM_018393.1 /DEF=Homo sapiens hypothetical protein FLJ11336 (FLJ11336), mRNA. /FEA=mRNA /GEN=FLJ11336 /PROD=hypothetical protein FLJ11336 /DB_XREF=gi:8922998 /UG=Hs.22383 hypothetical protein FLJ11336 /FL=gb:NM_018393.1
209731_at	gb:U79718.1 /DEF=Human endonuclease III homolog 1 (OCTS3) mRNA, complete cds. /FEA=mRNA /GEN=OCTS3 /PROD=endonuclease III homolog 1, hNTH1 /DB_XREF=gi:1800270 /UG=Hs.66196 nth (E.coli endonuclease III)-like 1 /FL=gb:BC000391.1 gb:BC003014.1 gb:U81285.1 gb:U79718.1 gb:AB001575.1 gb:NM_002528.3
212104_s_at	Consensus includes gb:N95026 /FEA=EST /DB_XREF=gi:1267308 /DB_XREF=est:zb45d12.s1 /CLONE=IMAGE:306551 /UG=Hs.5011 RNA binding motif protein 9
205801_s_at	gb:NM_015376.1 /DEF=Homo sapiens KIAA0846 protein (KIAA0846), mRNA. /FEA=mRNA /GEN=KIAA0846 /PROD=KIAA0846 protein /DB_XREF=gi:7662333 /UG=Hs.24024 KIAA0846 protein /FL=gb:AB020653.1
211064_at	gb:NM_015376.1
209502_s_at	gb:BC006408.1 /DEF=Homo sapiens, Similar to zinc finger protein 254, clone MGC:10544, mRNA, complete cds. /FEA=mRNA /PROD=Similar to zinc finger protein 254 /DB_XREF=gi:13623586 /FL=gb:BC006408.1
216158_at	gb:BC002495.1 /DEF=Homo sapiens, Similar to BAI1-associated protein 2, clone MGC:1836, mRNA, complete cds. /FEA=mRNA /PROD=Similar to BAI1-associated protein 2 /DB_XREF=gi:12803352 /UG=Hs.7936 BAI1-associated protein 2 /FL=gb:BC002495.1
210017_at	Consensus includes gb:AK024123.1 /DEF=Homo sapiens cDNA FLJ14061 fis, clone HEMBB1000749. /FEA=mRNA /DB_XREF=gi:10436427 /UG=Hs.306664 Homo sapiens cDNA FLJ14061 fis, clone HEMBB1000749
212014_x_at	Consensus includes gb:AF070528.1 /DEF=Homo sapiens clone 24631 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387883 /UG=Hs.180566 mucosa associated lymphoid tissue lymphoma translocation gene 1 /FL=gb:AB026118.1
209875_s_at	Consensus includes gb:AI493245 /FEA=EST /DB_XREF=gi:4394248 /DB_XREF=est:ti30d08.x1 /CLONE=IMAGE:2131983 /UG=Hs.169610 CD44 antigen (homing function and Indian blood group system)
207945_s_at	gb:M83248.1 /DEF=Human nephropontin mRNA, complete cds. /FEA=mRNA /GEN=nephropontin /PROD=nephropontin /DB_XREF=gi:189150 /UG=Hs.313 secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1) /FL=gb:M83248.1
	gb:NM_001893.1 /DEF=Homo sapiens casein kinase 1, delta (CSNK1D), mRNA. /FEA=mRNA /GEN=CSNK1D /PROD=casein kinase 1, delta /DB_XREF=gi:4503090 /UG=Hs.75852 casein kinase 1, delta /FL=gb:NM_001893.1
	gb:U29171.1

# Affy Probe\_ID

## Sequence Description

219874\_at gb:NM\_024628.1 /DEF=Homo sapiens hypothetical protein FLJ23188 (FLJ23188), mRNA. /FEA=mRNA /GEN=FLJ23188 /PROD=hypothetical protein FLJ23188 /DB\_XREF=gi:13375853 /UG=Hs.36793 hypothetical protein FLJ23188 /FL=gb:NM\_024628.1  
206715\_at gb:NM\_012252.1 /DEF=Homo sapiens transcription factor EC (TFEC), mRNA. /FEA=mRNA /GEN=TFEC /PROD=transcription factor EC /DB\_XREF=gi:6912701 /UG=Hs.113274 transcription factor EC /FL=gb:D43945.1 gb:NM\_012252.1  
200727\_s\_at Consensus includes gb:AA699583 /FEA=EST /DB\_XREF=gi:2703730 /DB\_XREF=est:zi42g07.s1 /CLONE=IMAGE:433500 /UG=Hs.42915 ARP2 (actin-related protein 2, yeast) homolog /FL=gb:AF006082.1 gb:NM\_005722.1  
220034\_at gb:NM\_007199.1 /DEF=Homo sapiens interleukin-1 receptor-associated kinase M (IRAK-M), mRNA. /FEA=mRNA /GEN=IRAK-M /PROD=interleukin-1 receptor-associated kinase M /DB\_XREF=gi:6005791 /UG=Hs.268552 interleukin-1 receptor-associated kinase M /FL=gb:AF113136.1 gb:NM\_007199.1  
212577\_at Consensus includes gb:AA868754 /FEA=EST /DB\_XREF=gi:2964199 /DB\_XREF=est:ak52e09.s1 /CLONE=IMAGE:1409608 /UG=Hs.8118 KIAA0650 protein  
205038\_at Consensus includes gb:BG540504 /FEA=EST /DB\_XREF=gi:13532737 /DB\_XREF=est:602569230F1 /CLONE=IMAGE:4693783 /UG=Hs.54452 zinc finger protein, subfamily 1A, 1 (Ikaros) /FL=gb:U40462.1 gb:NM\_006060.1  
219298\_at gb:NM\_024693.1 /DEF=Homo sapiens hypothetical protein FLJ20909 (FLJ20909), mRNA. /FEA=mRNA /GEN=FLJ20909 /PROD=hypothetical protein FLJ20909 /DB\_XREF=gi:13375972 /UG=Hs.98855 hypothetical protein FLJ20909 /FL=gb:NM\_024693.1  
219718\_at gb:NM\_018291.1 /DEF=Homo sapiens hypothetical protein FLJ10986 (FLJ10986), mRNA. /FEA=mRNA /GEN=FLJ10986 /PROD=hypothetical protein FLJ10986 /DB\_XREF=gi:8922805 /UG=Hs.273333 hypothetical protein FLJ10986 /FL=gb:BC000610.1 gb:NM\_018291.1  
204753\_s\_at Consensus includes gb:AI810712 /FEA=EST /DB\_XREF=gi:5397278 /DB\_XREF=est:tt86f08.x1 /CLONE=IMAGE:2248455 /UG=Hs.250692 hepatic leukemia factor /FL=gb:M95585.1 gb:NM\_002126.1  
202052\_s\_at gb:NM\_015577.1 /DEF=Homo sapiens novel retinal pigment epithelial gene (NORPEG), mRNA. /FEA=mRNA /GEN=NORPEG /PROD=DKFZP564G013 protein /DB\_XREF=gi:13470085 /UG=Hs.15165 novel retinal pigment epithelial gene /FL=gb:NM\_015577.1 gb:AF155135.1

# Affy Probe\_ID

## Sequence Description

215716\_s\_at Consensus includes gb:L14561 /DEF=Homo sapiens plasma membrane calcium ATPase isoform 1 (ATP2B1) gene, alternative splice products, partial cds /FEA=mRNA\_2 /DB\_XREF=gi:4165324 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1

211911\_x\_at gb:L07950.1 /DEF=Homo sapiens MHC class I HLA B71 mRNA, complete cds. /FEA=CDS /GEN=HLA-B /PROD=MHC HLA B71 /DB\_XREF=gi:307236 /FL=gb:L07950.1

203439\_s\_at gb:BC000658.1 /DEF=Homo sapiens, stanniocalcin 2, clone MGC:1881, mRNA, complete cds. /FEA=mRNA /PROD=stanniocalcin 2 /DB\_XREF=gi:12653744 /UG=Hs.155223 stanniocalcin 2 /FL=gb:BC000658.1 gb:AF055460.1 gb:AB012664.1 gb:AF098462.1 gb:AF031036.1 gb:NM\_003714.1

210136\_at Consensus includes gb:AW070431 /FEA=EST /DB\_XREF=gi:6025429 /DB\_XREF=est:xa11b04.x1 /CLONE=IMAGE:2567983 /UG=Hs.69547 myelin basic protein /FL=gb:L18865.1

201189\_s\_at gb:NM\_002224.1 /DEF=Homo sapiens inositol 1,4,5-triphosphate receptor, type 3 (ITPR3), mRNA. /FEA=mRNA /GEN=ITPR3 /PROD=inositol 1,4,5-triphosphate receptor, type 3 /DB\_XREF=gi:4504794 /UG=Hs.77515 inositol 1,4,5-triphosphate receptor, type 3 /FL=gb:D26351.1 gb:NM\_002224.1 gb:U01062.1

202014\_at gb:NM\_014330.2 /DEF=Homo sapiens growth arrest and DNA-damage-inducible 34 (GADD34), mRNA. /FEA=mRNA /GEN=GADD34 /PROD=growth arrest and DNA-damage-inducible 34 /DB\_XREF=gi:9790902 /UG=Hs.76556 growth arrest and DNA-damage-inducible 34 /FL=gb:BC003067.1 gb:U83981.1 gb:NM\_014330.2

200634\_at gb:NM\_005022.1 /DEF=Homo sapiens profilin 1 (PFN1), mRNA. /FEA=mRNA /GEN=PFN1 /PROD=profilin 1 /DB\_XREF=gi:4826897 /UG=Hs.75721 profilin 1 /FL=gb:BC002475.1 gb:J03191.1 gb:NM\_005022.1

203675\_at gb:NM\_005013.1 /DEF=Homo sapiens nucleobindin 2 (NUCB2), mRNA. /FEA=mRNA /GEN=NUCB2 /PROD=nucleobindin 2 /DB\_XREF=gi:4826869 /UG=Hs.3164 nucleobindin 2 /FL=gb:AF052642.1 gb:AF052643.1 gb:AF052644.1 gb:NM\_005013.1

221124\_s\_at gb:NM\_014588.2 /DEF=Homo sapiens visual system homeobox 1 (zebrafish) homolog (CHX10-like) (VSX1), mRNA. /FEA=mRNA /GEN=VSX1 /PROD=visual system homeobox 1 (zebrafish) homolog(CHX10-like)

214251\_s\_at /DB\_XREF=gi:11056037 /UG=Hs.274264 visual system homeobox 1 (zebrafish) homolog (CHX10-like) /FL=gb:AF176797.2 gb:NM\_014588.2 gb:AF251033.1

Consensus includes gb:AI337584 /FEA=EST /DB\_XREF=gi:4074511 /DB\_XREF=est:qw91a01.x1 /CLONE=IMAGE:1998408 /UG=Hs.301512 nuclear mitotic apparatus protein 1

Affy Probe_ID	Sequence Description
210868_s_at	gb:BC001305.1 /DEF=Homo sapiens, clone MGC:5487, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:5487) /DB_XREF=gi:12654918 /UG=Hs.211556 hypothetical protein MGC5487 /FL=gb:BC001305.1
201179_s_at	gb:J03005.1 /DEF=Human alternative guanine nucleotide-binding regulatory protein (G) alpha-inhibitory-subunit mRNA, complete cds. /FEA=mRNA /GEN=GNAI1 /DB_XREF=gi:183183 /UG=Hs.73799 guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3 /FL=gb:J03005.1 gb:J03198.1 gb:M27543.1 gb:J03238.1 gb:NM_006496.1
205289_at	Consensus includes gb:AA583044 /FEA=EST /DB_XREF=gi:2360404 /DB_XREF=est:nn80f11.s1 /CLONE=IMAGE:1090221 /UG=Hs.73853 bone morphogenetic protein 2 /FL=gb:NM_001200.1
217894_at	gb:NM_016121.1 /DEF=Homo sapiens NY-REN-45 antigen (LOC51133), mRNA. /FEA=mRNA /GEN=LOC51133 /PROD=NY-REN-45 antigen /DB_XREF=gi:7705836 /UG=Hs.239155 NY-REN-45 antigen /FL=gb:AF155110.1 gb:NM_016121.1
213486_at	Consensus includes gb:BF435376 /FEA=EST /DB_XREF=gi:11447664 /DB_XREF=est:nab38h02.x1 /CLONE=IMAGE:3268442 /UG=Hs.6421 hypothetical protein DKFZp761N09121
208955_at	gb:AB049113.1 /DEF=Homo sapiens DUT mRNA for dUTP pyrophosphatase, complete cds. /FEA=mRNA /GEN=DUT /PROD=dUTP pyrophosphatase /DB_XREF=gi:10257384 /UG=Hs.82113 dUTP pyrophosphatase /FL=gb:AB049113.1 gb:U31930.1 gb:U62891.1 gb:M89913.1 gb:NM_001948.1
202020_s_at	gb:NM_006055.1 /DEF=Homo sapiens LanC (bacterial lantibiotic synthetase component C)-like 1 (LANCL1), mRNA. /FEA=mRNA /GEN=LANCL1 /PROD=lanthionine synthetase C-like protein 1 /DB_XREF=gi:5174444 /UG=Hs.13351 LanC (bacterial lantibiotic synthetase component C)-like 1 /FL=gb:NM_006055.1
214209_s_at	Consensus includes gb:BE504895 /FEA=EST /DB_XREF=gi:9707303 /DB_XREF=est:h32b07.x1 /CLONE=IMAGE:3209653 /UG=Hs.6129 ATP-binding cassette, sub-family B (MDRTAP), member 9
202364_at	gb:NM_005962.1 /DEF=Homo sapiens MAX-interacting protein 1 (MXI1), mRNA. /FEA=mRNA /GEN=MXI1 /PROD=MAX-interacting protein 1 /DB_XREF=gi:5174596 /UG=Hs.118630 MAX-interacting protein 1 /FL=gb:L07648.1 gb:NM_005962.1 gb:D63940.1

# Affy Probe\_ID

## Sequence Description

207660\_at gb:NM\_004019.1 /DEF=Homo sapiens dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS270, DXS272 (DMD), transcript variant Dp40, mRNA. /FEA=mRNA /GEN=DMD /PROD=dystrophin Dp40 isoform /DB\_XREF=gi:5032304 /UG=Hs.169470 dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS270, DXS272 /FL=gb:NM\_004019.1

209627\_s\_at gb:AY008372.1 /DEF=Homo sapiens oxysterol binding protein-related protein 3 (ORP3) mRNA, complete cds. /FEA=mRNA /GEN=ORP3 /PROD=oxysterol binding protein-related protein 3 /DB\_XREF=gi:10880972 /UG=Hs.197955 KIAA0704 protein /FL=gb:AY008372.1

204733\_at gb:NM\_002774.1 /DEF=Homo sapiens kallikrein 6 (neurosin, zyme) (KLK6), mRNA. /FEA=mRNA /GEN=KLK6 /PROD=kallikrein 6 (neurosin, zyme) /DB\_XREF=gi:4506154 /UG=Hs.79361 kallikrein 6 (neurosin, zyme) /FL=gb:U62801.1 gb:D78203.1 gb:AF013988.1 gb:NM\_002774.1

201335\_s\_at gb:NM\_015313.1 /DEF=Homo sapiens KIAA0382 protein; leukemia-associated rho guanine nucleotide exchange factor (GEF) (ARHGEF12), mRNA. /FEA=mRNA /GEN=ARHGEF12 /PROD=KIAA0382 protein; leukemia-associated rhoguanine nucleotide exchange factor (GEF) /DB\_XREF=gi:7662087 /UG=Hs.6582 Rho guanine exchange factor (GEF) 12 /FL=gb:AF180681.1 gb:NM\_015313.1

204357\_s\_at gb:NM\_002314.2 /DEF=Homo sapiens LIM domain kinase 1 (LIMK1), transcript variant 1, mRNA. /FEA=mRNA /GEN=LIMK1 /PROD=LIM domain kinase 1 isoform 1 /DB\_XREF=gi:8051616 /UG=Hs.36566 LIM domain kinase 1 /FL=gb:D26309.1 gb:NM\_002314.2

209721\_s\_at gb:BC002857.1 /DEF=Homo sapiens, clone MGC:3442, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3442) /DB\_XREF=gi:12804008 /UG=Hs.46659 DKFZP586I2223 protein /FL=gb:BC002857.1 gb:BC001790.1 gb:NM\_015438.1 gb:BC004384.1

217998\_at Consensus includes gb:NM\_007350.1 /DEF=Homo sapiens pleckstrin homology-like domain, family A, member 1 (PHLDA1), mRNA. /FEA=mRNA /GEN=PHLDA1 /PROD=pleckstrin homology-like domain, family A, member 1 /DB\_XREF=gi:6679302 /UG=Hs.82101 pleckstrin homology-like domain, family A, member 1 /FL=gb:NM\_007350.1 gb:M87771.1 /DEF=Human secreted fibroblast growth factor receptor (K-sam-III) mRNA, complete cds. /FEA=mRNA /GEN=K-sam-III /PROD=fibroblast growth factor receptor /DB\_XREF=gi:186781 /UG=Hs.278581 fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) /FL=gb:NM\_022970.1 gb:M87771.1

# Affy Probe\_ID

## Sequence Description

215313_x_at	Consensus includes gb:AA573862 /FEA=EST /DB_XREF=gi:2348377 /DB_XREF=est:nk08f01.s1 /CLONE=IMAGE:1012921 /UG=Hs.181244 major histocompatibility complex, class I, A
220005_at	gb:NM_023914.1 /DEF=Homo sapiens G protein-coupled receptor 86 (GPR86), mRNA. /FEA=mRNA /GEN=GPR86 /PROD=G protein-coupled receptor 86 /DB_XREF=gi:13194202 /UG=Hs.13040 G protein-coupled receptor 86 /FL=gb:AF295368.1 gb:NM_023914.1 gb:AF178982.1 gb:AF345565.1
217916_s_at	gb:NM_016623.1 /DEF=Homo sapiens hypothetical protein (BM-009), mRNA. /FEA=mRNA /GEN=BM-009 /PROD=hypothetical protein /DB_XREF=gi:7705303 /UG=Hs.92918 hypothetical protein /FL=gb:BC003599.1 gb:AF208851.1 gb:NM_016623.1
216718_at	Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=gi:2589191 /UG=Hs.113261 Homo sapiens skin-specific protein (xp33) mRNA, partial cds
217794_at	gb:NM_018457.1 /DEF=Homo sapiens DKFZp564J157 protein (DKFZP564J157), mRNA. /FEA=mRNA /GEN=DKFZP564J157 /PROD=DKFZp564J157 protein /DB_XREF=gi:8922156 /UG=Hs.63042 DKFZp564J157 protein /FL=gb:AF217517.1 gb:NM_018457.1
204867_at	gb:NM_005258.2 /DEF=Homo sapiens GTP cyclohydrolase I feedback regulatory protein (GCHFR), mRNA. /FEA=mRNA /GEN=GCHFR /PROD=GTP cyclohydrolase I feedback regulatory protein /DB_XREF=gi:6382072 /UG=Hs.83081 GTP cyclohydrolase I feedback regulatory protein /FL=gb:NM_005258.2
218189_s_at	gb:NM_018946.2 /DEF=Homo sapiens N-acetylneuraminic acid phosphate synthase; sialic acid synthase (SAS), mRNA. /FEA=mRNA /GEN=SAS /PROD=N-acetylneuraminic acid phosphate synthase /DB_XREF=gi:12056472 /UG=Hs.274424 N-acetylneuraminic acid phosphate synthase; sialic acid synthase /FL=gb:NM_018946.2 gb:BC000008.1 gb:AF257466.1
201042_at	Consensus includes gb:AL031651 /DEF=Human DNA sequence from clone RP5-1054A22 on chromosome 20q11.22-12 Contains two isoforms of the gene for TGM2 (transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase), ESTs, STSs, GSSs and a CpG island /FEA=mRNA_1 /DB_XREF=gi:6065866 /UG=Hs.8265 transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase) /FL=gb:M55153.1 gb:NM_004613.1
218502_s_at	gb:NM_014112.1 /DEF=Homo sapiens trichorhinophalangeal syndrome I gene (TRPS1), mRNA. /FEA=mRNA /GEN=TRPS1 /PROD=trichorhinophalangeal syndrome I gene /DB_XREF=gi:7657658 /UG=Hs.26102 trichorhinophalangeal syndrome I /FL=gb:AF183810.1 gb:NM_014112.1

Affy Probe_ID	Sequence Description
201906_s_at	gb:NM_005808.1 /DEF=Homo sapiens HYA22 protein (HYA22), mRNA. /FEA=mRNA /GEN=HYA22 /PROD=HYA22 protein /DB_XREF=gi:5031774 /UG=Hs.147189 HYA22 protein /FL=gb:D88153.1 gb:NM_005808.1 gb:D21254.1 /DEF=Human mRNA for OB-cadherin-1, complete cds. /FEA=mRNA /GEN=osf-4 /PROD=OB-cadherin-1 /DB_XREF=gi:575577 /UG=Hs.75929 cadherin 11, type 2, OB-cadherin (osteoblast) /FL=gb:NM_001797.1 gb:L34056.1 gb:D21254.1
200825_s_at	gb:NM_006389.2 /DEF=Homo sapiens oxygen regulated protein (150kD) (ORP150), mRNA. /FEA=mRNA /GEN=ORP150 /PROD=oxygen regulated protein precursor /DB_XREF=gi:13699861 /UG=Hs.277704 oxygen regulated protein (150kD) /FL=gb:NM_006389.2 gb:U65785.1
212068_s_at	Consensus includes gb:AB011087.1 /DEF=Homo sapiens mRNA for KIAA0515 protein, partial cds. /FEA=mRNA /GEN=KIAA0515 /PROD=KIAA0515 protein /DB_XREF=gi:3043553 /UG=Hs.108945 KIAA0515 protein
221485_at	Consensus includes gb:AL035683 /DEF=Human DNA sequence from clone RP5-1063B2 on chromosome 20q13.1-13.2. Contains the 3 part of the gene for Beta-1,4-galactosyltransferase, ESTs, STSs and GSSs /FEA=mRNA /DB_XREF=gi:7288039 /UG=Hs.107526 UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5 /FL=gb:AB004550.1 gb:AF038663.1 gb:NM_004776.1
209834_at	gb:AB017915.1 /DEF=Homo sapiens mRNA for chondroitin 6-sulfotransferase, complete cds. /FEA=mRNA /PROD=chondroitin 6-sulfotransferase /DB_XREF=gi:4115403 /UG=Hs.158304 carbohydrate (chondroitin 6keratan) sulfotransferase 3 /FL=gb:AB017915.1
219540_at	Consensus includes gb:AU150728 /FEA=EST /DB_XREF=gi:11012249 /DB_XREF=est:AU150728 /CLONE=NT2RP2003470 /UG=Hs.145498 zinc finger protein 267 /FL=gb:NM_003414.2 gb:AF220492.1
217473_x_at	Consensus includes gb:AF229163 /DEF=Homo sapiens natural resistance-associated macrophage protein 1 (SLC11A1) gene, complete cds, alternatively spliced; and nuclear LIM interactor-interacting factor (NLI-IF) gene, complete cds /FEA=mRNA_3 /DB_XREF=gi:10257408 /UG=Hs.182611 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1
220751_s_at	gb:NM_016348.1 /DEF=Homo sapiens chromosome 5 open reading frame 4 (C5ORF4), mRNA. /FEA=mRNA /GEN=C5ORF4 /PROD=putative tumor suppressor /DB_XREF=gi:7705942 /UG=Hs.10235 chromosome 5 open reading frame 4 /FL=gb:AF159165.1 gb:NM_016348.1
213523_at	Consensus includes gb:AI671049 /FEA=EST /DB_XREF=gi:4850780 /DB_XREF=est:tz17d11.x1 /CLONE=IMAGE:2288853 /UG=Hs.9700 cyclin E1

**Affy Probe ID**

**Sequence Description**

204922_at	gb:NM_024650.1 /DEF=Homo sapiens hypothetical protein FLJ22531 (FLJ22531), mRNA. /FEA=mRNA /GEN=FLJ22531 /PROD=hypothetical protein FLJ22531 /DB_XREF=gi:13375894 /UG=Hs.55613 hypothetical protein FLJ22531 /FL=gb:NM_024650.1
212364_at	Consensus includes gb:BF432550 /FEA=EST /DB_XREF=gi:11444700 /DB_XREF=est:nac56e10.x1 /CLONE=IMAGE:3406555 /UG=Hs.121576 Homo sapiens cDNA FLJ20153 fis, clone COL08656, highly similar to AJ001381 Homo sapiens incomplete cDNA for a mutated allele
202724_s_at	gb:NM_002015.2 /DEF=Homo sapiens forkhead box O1A (rhabdomyosarcoma) (FOXO1A), mRNA. /FEA=mRNA /GEN=FOXO1A /PROD=forkhead box O1A /DB_XREF=gi:9257221 /UG=Hs.170133 forkhead box O1A (rhabdomyosarcoma) /FL=gb:AF032885.1 gb:U02310.1 gb:NM_002015.2
212636_at	Consensus includes gb:AL031781 /DEF=Human DNA sequence from clone 51J12 on chromosome 6q26-27. Contains the 3 part of the alternatively spliced gene for the human orthologs of mouse QKI-7 and QKI-7B (KH Domain RNA Binding proteins) and zebrafish ZKQ-1 (Quaking protein homolog). Con... /FEA=mRNA_2 /DB_XREF=gi:4038570 /UG=Hs.15020 homolog of mouse quaking QKI (KH domain RNA binding protein) /FL=gb:AF142417.1
215078_at	Consensus includes gb:AL050388.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M2422 (from clone DKFZp564M2422); partial cds. /FEA=mRNA /GEN=DKFZp564M2422 /PROD=hypothetical protein /DB_XREF=gi:4914612 /UG=Hs.306320 Homo sapiens mRNA; cDNA DKFZp564M2422 (from clone DKFZp564M2422); partial cds
203697_at	gb:U91903.1 /DEF=Human Fritz mRNA, complete cds. /FEA=mRNA /PROD=Fritz /DB_XREF=gi:1917006 /UG=Hs.153684 frizzled-related protein /FL=gb:U24163.1 gb:U68057.1 gb:U91903.1 gb:NM_001463.1
203620_s_at	gb:NM_014824.1 /DEF=Homo sapiens KIAA0769 gene product (KIAA0769), mRNA. /FEA=mRNA /GEN=KIAA0769 /PROD=KIAA0769 gene product /DB_XREF=gi:7662295 /UG=Hs.19056 KIAA0769 gene product /FL=gb:AB018312.1 gb:NM_014824.1
201502_s_at	Consensus includes gb:AI078167 /FEA=EST /DB_XREF=gi:3412575 /DB_XREF=est:oz30d08.x1 /CLONE=IMAGE:1676847 /UG=Hs.81328 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha /FL=gb:NM_020529.1 gb:BC002601.1 gb:BC004983.1 gb:M69043.1
205568_at	gb:NM_020980.2 /DEF=Homo sapiens aquaporin 9 (AQP9), mRNA. /FEA=mRNA /GEN=AQP9 /PROD=aquaporin 9 /DB_XREF=gi:11038652 /UG=Hs.104624 aquaporin 9 /FL=gb:NM_020980.2 gb:AB008775.1 gb:AF016495.1
210118_s_at	gb:M15329.1 /DEF=Human interleukin 1-alpha (IL1A) mRNA, complete cds. /FEA=mRNA /GEN=IL1A /PROD=interleukin 1-alpha /DB_XREF=gi:186277 /UG=Hs.1722 interleukin 1, alpha /FL=gb:M15329.1



# Affy Probe ID

## Sequence Description

219664_s_at	gb:NM_020664.1 /DEF=Homo sapiens 2,4-dienoyl CoA reductase 2, peroxisomal (DEC2), mRNA. /FEA=mRNA /GEN=DEC2 /PROD=peroxisomal 2,4-dienoyl-CoA reductase /DB_XREF=gi:10190703 /UG=Hs.15898 2,4-dienoyl CoA reductase 2, peroxisomal /FL=gb:NM_020664.1
216215_s_at	Consensus includes gb:AL049748 /DEF=Human DNA sequence from clone RP1-41P2 on chromosome 22 Contains the 3 part of the RBM9 gene for RNA binding motif protein 9 and the 3 part of the gene for a novel protein similar to part of APOL (apolipoprotein L) and TNF-inducible protein CG12... /FEA=mRNA_2 /DB_XREF=gi:6572235 /UG=Hs.5011 RNA binding motif protein 9
202450_s_at	gb:NM_000396.1 /DEF=Homo sapiens cathepsin K (pseudosostosis) (CTSK), mRNA. /FEA=mRNA /GEN=CTSK /PROD=cathepsin K (pseudosostosis) /DB_XREF=gi:4503150 /UG=Hs.83942 cathepsin K (pseudosostosis) /FL=gb:NM_000396.1 gb:U13665.1
219869_s_at	gb:NM_022154.1 /DEF=Homo sapiens up-regulated by BCG-CWS (LOC64116), mRNA. /FEA=mRNA /GEN=LOC64116 /PROD=up-regulated by BCG-CWS /DB_XREF=gi:11545899 /UG=Hs.284205 up-regulated by BCG-CWS /FL=gb:NM_022154.1
200736_s_at	gb:NM_000581.1 /DEF=Homo sapiens glutathione peroxidase 1 (GPX1), mRNA. /FEA=mRNA /GEN=GPX1 /PROD=glutathione peroxidase 1 /DB_XREF=gi:10834975 /UG=Hs.76686 glutathione peroxidase 1 /FL=gb:NM_000581.1 gb:BC000742.1 gb:M21304.1
204646_at	gb:NM_000110.2 /DEF=Homo sapiens dihydropyrimidine dehydrogenase (DPYD), mRNA. /FEA=mRNA /GEN=DPYD /PROD=dihydropyrimidine dehydrogenase /DB_XREF=gi:4557874 /UG=Hs.1602 dihydropyrimidine dehydrogenase /FL=gb:U20938.1 gb:NM_000110.2 gb:U09178.1 gb:AB003063.1
204029_at	gb:NM_001408.1 /DEF=Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 2, flamingo (Drosophila) homolog (CELSR2), mRNA. /FEA=mRNA /GEN=CELSR2 /PROD=cadherin EGF LAG seven-pass G-type receptor 2 /DB_XREF=gi:13325063 /UG=Hs.57652 cadherin, EGF LAG seven-pass G-type receptor 2, flamingo (Drosophila) homolog /FL=gb:NM_001408.1
200871_s_at	gb:NM_002778.1 /DEF=Homo sapiens prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy) (PSAP), mRNA. /FEA=mRNA /GEN=PSAP /PROD=prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy) /DB_XREF=gi:11386146 /UG=Hs.78575 prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy) /FL=gb:NM_002778.1 gb:BC004275.1 gb:J03077.1 gb:D00422.1 gb:M60255.1 gb:M32221.1 gb:M60257.1 gb:M60258.1 gb:M81355.1

**Affy Probe ID**

**Sequence Description**

211670_x_at	gb:S82471.1 /DEF=Homo sapiens Kruppel-associated box containing gene product SSX3 (SSX3) mRNA, complete cds. /FEA=mRNA /GEN=SSX3 /PROD=Kruppel-associated box containing gene productSSX3 /DB_XREF=gi:1699271 /FL=gb:S82471.1
218613_at	gb:NM_018422.1 /DEF=Homo sapiens hypothetical protein DKFZp761K1423 (DKFZp761K1423), mRNA. /FEA=mRNA /GEN=DKFZp761K1423 /PROD=hypothetical protein DKFZp761K1423 /DB_XREF=gi:8922171 /UG=Hs.236438 hypothetical protein DKFZp761K1423 /FL=gb:NM_018422.1
200629_at	gb:NM_004184.2 /DEF=Homo sapiens tryptophanyl-tRNA synthetase (WARS), mRNA. /FEA=mRNA /GEN=WARS /PROD=tryptophanyl-tRNA synthetase /DB_XREF=gi:7710155 /UG=Hs.82030 tryptophanyl-tRNA synthetase /FL=gb:M77804.1 gb:M61715.1 gb:NM_004184.2
202307_s_at	gb:NM_000593.2 /DEF=Homo sapiens ATP-binding cassette, sub-family B (MDRTAP), member 2 (ABCB2), mRNA. /FEA=mRNA /GEN=ABCB2 /PROD=ATP-binding cassette, sub-family B, member 2 /DB_XREF=gi:9665247 /UG=Hs.158164 ATP-binding cassette, sub-family B (MDRTAP), member 2 /FL=gb:L21204.1 gb:L21205.1 gb:L21206.1 gb:L21207.1 gb:L21208.1 gb:NM_000593.2
203175_at	gb:NM_001665.1 /DEF=Homo sapiens ras homolog gene family, member G (rho G) (ARHG), mRNA. /FEA=mRNA /GEN=ARHG /PROD=ras homolog gene family, member G (rho G) /DB_XREF=gi:4502218 /UG=Hs.75082 ras homolog gene family, member G (rho G) /FL=gb:NM_001665.1
203382_s_at	gb:NM_000041.1 /DEF=Homo sapiens apolipoprotein E (APOE), mRNA. /FEA=mRNA /GEN=APOE /PROD=apolipoprotein E /DB_XREF=gi:4557324 /UG=Hs.169401 apolipoprotein E /FL=gb:BC003557.1 gb:M12529.1 gb:K00396.1 gb:NM_000041.1
209281_s_at	gb:M95541.1 /DEF=Homo sapiens adenosine triphosphatase mRNA, complete cds. /FEA=mRNA /PROD=adenosine triphosphatase /DB_XREF=gi:184269 /UG=Hs.78546 ATPase, Ca++ transporting, plasma membrane 1 /FL=gb:M95541.1 gb:NM_001682.1 gb:J04027.1
201313_at	gb:NM_001975.1 /DEF=Homo sapiens enolase 2, (gamma, neuronal) (ENO2), mRNA. /FEA=mRNA /GEN=ENO2 /PROD=enolase 2, (gamma, neuronal) /DB_XREF=gi:5803010 /UG=Hs.146580 enolase 2, (gamma, neuronal) /FL=gb:BC002745.1 gb:NM_001975.1 gb:M22349.1
208012_x_at	gb:NM_004509.1 /DEF=Homo sapiens interferon-induced protein 41, 30kD (IFI41), mRNA. /FEA=mRNA /GEN=IFI41 /PROD=interferon-induced protein 41, 30kD /DB_XREF=gi:4758585 /UG=Hs.241510 interferon-induced protein 41, 30kD /FL=gb:L22342.1 gb:NM_004509.1

# Affy Probe\_ID

## Sequence Description

203028\_s\_at gb:NM\_000101.1 /DEF=Homo sapiens cytochrome b-245, alpha polypeptide (CYBA), mRNA. /FEA=mRNA /GEN=CYBA /PROD=flavocytochrome b-558 alpha polypeptide /DB\_XREF=gi:4557504 /UG=Hs.68877 cytochrome b-245, alpha polypeptide /FL=gb:M21186.1 gb:NM\_000101.1

207277\_at gb:AF290886.1 /DEF=Homo sapiens DC-SIGN mRNA, complete cds. /FEA=mRNA /PROD=DC-SIGN /DB\_XREF=gi:13383467 /UG=Hs.278694 CD209 antigen /FL=gb:NM\_021155.1 gb:AF290886.1 gb:M98457.1

212235\_at Consensus includes gb:AL575403 /FEA=EST /DB\_XREF=gi:12936533 /DB\_XREF=est:AL575403 /CLONE=CS0DI060YL02 (3 prime) /UG=Hs.301685 KIAA0620 protein

209537\_at gb:AF000416.1 /DEF=Homo sapiens EXT-like protein 2 (EXTL2) mRNA, complete cds. /FEA=mRNA /GEN=EXTL2 /PROD=EXT-like protein 2 /DB\_XREF=gi:2895061 /UG=Hs.61152 exostoses (multiple)-like 2 /FL=gb:AB009284.1 gb:AF000416.1

213935\_at Consensus includes gb:AF007132.1 /DEF=Homo sapiens clone 23551 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:2852606 /UG=Hs.184019 Homo sapiens clone 23551 mRNA sequence

220615\_s\_at gb:NM\_018099.1 /DEF=Homo sapiens hypothetical protein FLJ10462 (FLJ10462), mRNA. /FEA=mRNA /GEN=FLJ10462 /PROD=hypothetical protein FLJ10462 /DB\_XREF=gi:8922433 /UG=Hs.100895 hypothetical protein FLJ10462 /FL=gb:AL136843.1 gb:NM\_018099.1

221527\_s\_at gb:AF196185.1 /DEF=Homo sapiens atypical PKC isotype-specific interacting protein long variant mRNA, complete cds. /FEA=mRNA /PROD=atypical PKC isotype-specific interacting protein long variant /DB\_XREF=gi:13491609 /UG=Hs.72249 three-PDZ containing protein similar to C. elegans PAR3 (partitioning defect) /FL=gb:AF196185.1

209500\_x\_at gb:AF114012.1 /DEF=Homo sapiens tumor necrosis factor-related death ligand-1 beta mRNA, complete cds. /FEA=mRNA /PROD=tumor necrosis factor-related death ligand-1 beta /DB\_XREF=gi:7328555 /UG=Hs.54673 tumor necrosis factor (ligand) superfamily, member 13 /FL=gb:AF114012.1

221833\_at Consensus includes gb:AI971258 /FEA=EST /DB\_XREF=gi:5768084 /DB\_XREF=est:wr27b08.x1 /CLONE=IMAGE:248887 /UG=Hs.295923 seven in absentia (Drosophila) homolog 1

209189\_at gb:BC004490.1 /DEF=Homo sapiens, v-fos FBJ murine osteosarcoma viral oncogene homolog, clone MGC:11074, mRNA, complete cds. /FEA=mRNA /PROD=v-fos FBJ murine osteosarcoma viral oncogene homolog /DB\_XREF=gi:13325363 /UG=Hs.25647 v-fos FBJ murine osteosarcoma viral oncogene homolog /FL=gb:BC004490.1 gb:NM\_005252.2

**Affy Probe ID**

**Sequence Description**

201780\_s\_at

gb:NM\_007282.1 /DEF=Homo sapiens ring finger protein 13 (RNF13), mRNA. /FEA=mRNA /GEN=RNF13 /PROD=ring finger protein 13 /DB\_XREF=gi:6005863 /UG=Hs.6900 ring finger protein 13 /FL=gb:AF037204.1 gb:AF070558.1 gb:NM\_007282.1

203019\_x\_at

gb:NM\_014021.1 /DEF=Homo sapiens KIAA0923 protein (KIAA0923), mRNA. /FEA=mRNA /GEN=KIAA0923 /PROD=KIAA0923 protein /DB\_XREF=gi:7662381 /UG=Hs.22587 KIAA0923 protein /FL=gb:AB023140.1 gb:NM\_014021.1

212533\_at

Consensus includes gb:X62048.1 /DEF=H.sapiens Weel hu gene. /FEA=mRNA /GEN=Weel Hu /DB\_XREF=gi:499071 /UG=Hs.75188 weel+ (S. pombe) homolog /FL=gb:NM\_003390.1 gb:U10564.1

205107\_s\_at

gb:NM\_005227.1 /DEF=Homo sapiens ephrin-A4 (EFNA4), mRNA. /FEA=mRNA /GEN=EFNA4 /PROD=ephrin-A4 /DB\_XREF=gi:4885196 /UG=Hs.42331 ephrin-A4 /FL=gb:NM\_005227.1 gb:U14188.1

218257\_s\_at

gb:NM\_020120.1 /DEF=Homo sapiens UDP-glucose:glycoprotein glucosyltransferase 1 (HUGT1), mRNA. /FEA=mRNA /GEN=HUGT1 /PROD=UDP-glucose:glycoprotein glucosyltransferase 1 /DB\_XREF=gi:9910279

203923\_s\_at

UG=Hs.105794 UDP-glucose:glycoprotein glucosyltransferase 1 /FL=gb:AF227905.1 gb:NM\_020120.1 gb:NM\_000397.2 /DEF=Homo sapiens cytochrome b-245, beta polypeptide (chronic granulomatous disease) (CYBB), mRNA. /FEA=mRNA /GEN=CYBB /PROD=cytochrome b-245, beta polypeptide (chronic granulomatous disease) /DB\_XREF=gi:6996020 /UG=Hs.88974 cytochrome b-245, beta polypeptide (chronic granulomatous disease) /FL=gb:NM\_000397.2

204698\_at

gb:NM\_002201.2 /DEF=Homo sapiens interferon stimulated gene (20kD) (ISG20), mRNA. /FEA=mRNA /GEN=ISG20 /PROD=interferon stimulated gene (20kD) /DB\_XREF=gi:6857799 /UG=Hs.183487 interferon stimulated gene (20kD) /FL=gb:U88964.1 gb:NM\_002201.2

202587\_s\_at

gb:BC001116.1 /DEF=Homo sapiens, adenylate kinase 1, clone MGC:1808, mRNA, complete cds. /FEA=mRNA /PROD=adenylate kinase 1 /DB\_XREF=gi:12654562 /UG=Hs.76240 adenylate kinase 1 /FL=gb:BC001116.1 gb:NM\_000476.1 gb:AB021871.1

205082\_s\_at

gb:AB046692.1 /DEF=Homo sapiens hAO mRNA for aldehyde oxidase, complete cds. /FEA=mRNA /GEN=hAO /PROD=aldehyde oxidase /DB\_XREF=gi:13516378 /UG=Hs.174151 aldehyde oxidase 1 /FL=gb:AB046692.1 gb:L11005.1 gb:NM\_001159.2

207016\_s\_at

gb:AB015228.1 /DEF=Homo sapiens mRNA for RALDH2-T, complete cds. /FEA=mRNA /GEN=RALDH2 /PROD=RALDH2-T /DB\_XREF=gi:3970845 /UG=Hs.95197 aldehyde dehydrogenase 1 family, member A2 /FL=gb:NM\_003888.1 gb:AB015226.1 gb:AB015227.1 gb:AB015228.1

Affy Probe_ID	Sequence Description
206978_at	gb:NM_000647.2 /DEF=Homo sapiens chemokine (C-C motif) receptor 2 (CCR2), mRNA. /FEA=mRNA /GEN=CCR2 /PROD=chemokine (C-C motif) receptor 2 /DB_XREF=gi:4827072 /UG=Hs.395 chemokine (C-C motif) receptor 2 /FL=gb:U03882.1 gb:NM_000647.2
205176_s_at	gb:NM_014288.1 /DEF=Homo sapiens integrin beta 3 binding protein (beta3-endonexin) (ITGB3BP), mRNA. /FEA=mRNA /GEN=ITGB3BP /PROD=integrin beta 3 binding protein(beta3-endonexin) /DB_XREF=gi:7657205 /UG=Hs.82084 integrin beta 3 binding protein (beta3-endonexin) /FL=gb:BC005301.1 gb:AF175306.1 gb:NM_014288.1
202887_s_at	gb:NM_019058.1 /DEF=Homo sapiens hypothetical protein (FLJ20500), mRNA. /FEA=mRNA /GEN=FLJ20500 /PROD=hypothetical protein /DB_XREF=gi:9506686 /UG=Hs.111244 hypothetical protein /FL=gb:AL136668.1 gb:NM_019058.1
214452_at	Consensus includes gb:NM_005504.1 /DEF=Homo sapiens branched chain aminotransferase 1, cytosolic (BCAT1), mRNA. /FEA=CDS /GEN=BCAT1 /PROD=branched chain aminotransferase 1, cytosolic /DB_XREF=gi:5031606 /UG=Hs.157205 branched chain aminotransferase 1, cytosolic /FL=gb:U21551.1 gb:NM_005504.1
209108_at	gb:AF053453.1 /DEF=Homo sapiens tetraspan TM4SF (TSPAN-6) mRNA, complete cds. /FEA=mRNA /GEN=TSPAN-6 /PROD=tetraspan TM4SF /DB_XREF=gi:2995860 /UG=Hs.121068 transmembrane 4 superfamily member 6 /FL=gb:AF043906.1 gb:AF053453.1 gb:U84895.1 gb:NM_003270.1 gb:AF133426.1
213385_at	Consensus includes gb:AK026415.1 /DEF=Homo sapiens cDNA: FLJ22762 fis, clone KAI0960, highly similar to HUMB2CHIM Homo sapiens beta2-chimaerin mRNA. /FEA=mRNA /DB_XREF=gi:10439275 /UG=Hs.286055 chimerin (chimaerin) 2
204755_x_at	gb:M95585.1 /DEF=Human hepatic leukemia factor (HLF) mRNA, complete cds. /FEA=mRNA /GEN=HLF /PROD=hepatic leukemia factor /DB_XREF=gi:184223 /UG=Hs.250692 hepatic leukemia factor /FL=gb:M95585.1 gb:NM_002126.1
212775_at	Consensus includes gb:AI978623 /FEA=EST /DB_XREF=gi:5803653 /DB_XREF=est:wr57d03.x1 /CLONE=IMAGE:2491781 /UG=Hs.6654 KIAA0657 protein
213482_at	Consensus includes gb:BF593175 /FEA=EST /DB_XREF=gi:11685499 /DB_XREF=est:7o57g10.x1 /CLONE=IMAGE:3578322 /UG=Hs.7022 dedicator of cyto-kinesis 3
205191_at	gb:NM_006915.1 /DEF=Homo sapiens retinitis pigmentosa 2 (X-linked recessive) (RP2), mRNA. /FEA=mRNA /GEN=RP2 /PROD=XRP2 protein /DB_XREF=gi:5902059 /UG=Hs.44766 retinitis pigmentosa 2 (X-linked recessive) /FL=gb:NM_006915.1

# Affy Probe\_ID

## Sequence Description

217867_x_at	gb:NM_012105.1 /DEF=Homo sapiens beta-site APP-cleaving enzyme 2 (BACE2), mRNA. /FEA=mRNA /GEN=BACE2 /PROD=beta-site APP-cleaving enzyme 2 /DB_XREF=gi:6912263 /UG=Hs.271411 beta-site APP-cleaving enzyme 2 /FL=gb:AF117892.1 gb:AF050171.1 gb:AF200192.1 gb:AF200342.1 gb:AF204944.1 gb:AF178532.1 gb:NM_012105.1
216464_x_at	Consensus includes gb:AF118265 /DEF=Homo sapiens orphan G protein-coupled receptor GPR44 (GPR44) gene, complete cds /FEA=CDS /DB_XREF=gi:4455060 /UG=Hs.299567 G protein-coupled receptor 44
215239_x_at	Consensus includes gb:AU132789 /FEA=EST /DB_XREF=gi:10993328 /DB_XREF=est:AU132789 /CLONE=NT2RP4000545 /UG=Hs.89732 zinc finger protein 273
205554_s_at	gb:NM_004944.1 /DEF=Homo sapiens deoxyribonuclease I-like 3 (DNASE1L3), mRNA. /FEA=mRNA /GEN=DNASE1L3 /PROD=deoxyribonuclease I-like 3 /DB_XREF=gi:4826697 /UG=Hs.88646 deoxyribonuclease I-like 3 /FL=gb:U56814.1 gb:AF047354.1 gb:NM_004944.1
218627_at	gb:NM_018370.1 /DEF=Homo sapiens hypothetical protein FLJ11259 (FLJ11259), mRNA. /FEA=mRNA /GEN=FLJ11259 /PROD=hypothetical protein FLJ11259 /DB_XREF=gi:8922957 /UG=Hs.184465 hypothetical protein FLJ11259 /FL=gb:NM_018370.1
217901_at	Consensus includes gb:BF031829 /FEA=EST /DB_XREF=gi:10739541 /DB_XREF=est:601557983F1 /CLONE=IMAGE:3827709 /UG=Hs.2631 desmoglein 2 /FL=gb:NM_001943.1
209073_s_at	gb:AF015040.1 /DEF=Homo sapiens NUMB protein (NUMB) mRNA, complete cds. /FEA=mRNA /GEN=NUMB /PROD=NUMB protein /DB_XREF=gi:4102704 /UG=Hs.78890 numb (Drosophila) homolog /FL=gb:AF015040.1 gb:AF108092.1 gb:AF171939.1
204950_at	gb:NM_014959.1 /DEF=Homo sapiens KIAA0955 protein (KIAA0955), mRNA. /FEA=mRNA /GEN=KIAA0955 /PROD=KIAA0955 protein /DB_XREF=gi:7662403 /UG=Hs.10031 KIAA0955 protein /FL=gb:AF322184.1 gb:AF331519.1 gb:AY026322.1 gb:AB023172.1 gb:NM_014959.1
203340_s_at	gb:NM_003705.1 /DEF=Homo sapiens solute carrier family 25 (mitochondrial carrier, Aralar), member 12 (SLC25A12), mRNA. /FEA=mRNA /GEN=SLC25A12 /PROD=solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /DB_XREF=gi:4507006 /UG=Hs.179866 solute carrier family 25 (mitochondrial carrier, Aralar), member 12 /FL=gb:NM_003705.1
211423_s_at	gb:D85181.1 /DEF=Homo sapiens mRNA for fungal sterol-C5-desaturase homolog, complete cds. /FEA=mRNA /PROD=fungal sterol-C5-desaturase homolog /DB_XREF=gi:1906795 /UG=Hs.288031 sterol-C5-desaturase (fungal ERG3, delta-5-desaturase)-like /FL=gb:D85181.1

Affy Probe_ID	Sequence Description
212561_at	Consensus includes gb:AA349595 /FEA=EST /DB_XREF=gi:2001934 /DB_XREF=est:EST56429 /UG=Hs.26797 KIAA1091 protein
210660_at	gb:AF025529.1 /DEF=Homo sapiens leucocyte immunoglobulin-like receptor-6b (LIR-6) mRNA, complete cds. /FEA=mRNA /GEN=LIR-6 /PROD=leucocyte immunoglobulin-like receptor-6b /DB_XREF=gi:2653864 /UG=Hs.166156 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1 /FL=gb:AF025529.1
210313_at	gb:AF041261.1 /DEF=Homo sapiens immunoglobulin-like transcript 7 mRNA, complete cds. /FEA=mRNA /PROD=immunoglobulin-like transcript 7 /DB_XREF=gi:4104890 /UG=Hs.48647 immunoglobulin-like transcript 7 /FL=gb:AF041261.1
217807_s_at	gb:NM_015710.1 /DEF=Homo sapiens glioma tumor suppressor candidate region gene 2 (GLTSCR2), mRNA. /FEA=mRNA /GEN=GLTSCR2 /PROD=glioma tumor suppressor candidate region gene 2 /DB_XREF=gi:7657129 /UG=Hs.2237 glioma tumor suppressor candidate region gene 2 /FL=gb:AF182076.1 gb:NM_015710.1
212930_at	Consensus includes gb:AW576457 /FEA=EST /DB_XREF=gi:7247996 /DB_XREF=est:UI-HF-BR0p-aji-a-01-0-UI.s1 /CLONE=IMAGE:3074568 /UG=Hs.20952 Homo sapiens clone 24411 mRNA sequence
213212_x_at	Consensus includes gb:AI632181 /FEA=EST /DB_XREF=gi:4683511 /DB_XREF=est:ts85h05.x1 /CLONE=IMAGE:2238105 /UG=Hs.182982 golgin-67
203567_s_at	Consensus includes gb:AU157590 /FEA=EST /DB_XREF=gi:11019111 /DB_XREF=est:AU157590 /CLONE=PLACE1008438 /UG=Hs.59545 ring finger protein 15 /FL=gb:U90547.1 gb:NM_006355.1
219342_at	gb:NM_022900.1 /DEF=Homo sapiens hypothetical protein FLJ21213 (FLJ21213), mRNA. /FEA=mRNA /GEN=FLJ21213 /PROD=hypothetical protein FLJ21213 /DB_XREF=gi:12597638 /UG=Hs.128003 hypothetical protein FLJ21213 /FL=gb:NM_022900.1
212702_s_at	Consensus includes gb:N45111 /FEA=EST /DB_XREF=gi:1186277 /DB_XREF=est:yz12f12.s1 /CLONE=IMAGE:282863 /UG=Hs.330988 Homo sapiens, Similar to Bicaudal D (Drosophila) homolog 1, clone IMAGE:3622452, mRNA, partial cds
209340_at	gb:S73498.1 /DEF=Homo sapiens AgX-1 antigen mRNA, complete cds. /FEA=mRNA /PROD=AgX-1 antigen /DB_XREF=gi:688010 /UG=Hs.21293 UDP-N-acteylglucosamine pyrophosphorylase 1 /FL=gb:AB011004.1 gb:NM_003115.1 gb:S73498.1

Affy Probe_ID	Sequence Description
202943_s_at	gb:M38083.1 /DEF=Human alpha-N-acetylgalactosaminidase mRNA, complete cds. /FEA=mRNA /PROD=alpha-N-acetylgalactosaminidase /DB_XREF=gi:189054 /UG=Hs.75372 N-acetylgalactosaminidase, alpha- /FL=gb:BC000095.1 gb:M62783.1 gb:M38083.1 gb:NM_000262.1
202894_at	gb:NM_004444.1 /DEF=Homo sapiens EphB4 (EPHB4) mRNA. /FEA=mRNA /GEN=HTK /PROD=EphB4 /DB_XREF=gi:4758289 /UG=Hs.155227 EphB4 /FL=gb:NM_004444.1 gb:U07695.1
207670_at	gb:NM_002283.1 /DEF=Homo sapiens keratin, hair, basic, 5 (KRTHB5), mRNA. /FEA=mRNA /GEN=KRTHB5 /PROD=keratin, hair, basic, 5 /DB_XREF=gi:4504934 /UG=Hs.182507 keratin, hair, basic, 5 /FL=gb:NM_002283.1
202375_at	gb:NM_014822.1 /DEF=Homo sapiens SEC24 (S. cerevisiae) related gene family, member D (SEC24D), mRNA. /FEA=mRNA /GEN=SEC24D /PROD=SEC24 (S. cerevisiae) related gene family, member D /DB_XREF=gi:7662658 /UG=Hs.19822 SEC24 (S. cerevisiae) related gene family, member D /FL=gb:AB018298.1 gb:AF130464.2 gb:NM_014822.1
39318_at	Cluster Incl. X82240:H.sapiens mRNA for Tcell leukemia/lymphoma 1 /cds=(45,389) /gb=X82240 /gi=624960 /ug=Hs.2484 /len=1312
221765_at	Consensus includes gb:AI378044 /FEA=EST /DB_XREF=gi:4187897 /DB_XREF=est:te67h09.x1 /CLONE=IMAGE:2091809 /UG=Hs.23703 ESTs
202746_at	Consensus includes gb:AL021786 /DEF=Human DNA sequence from PAC 696H22 on chromosome Xq21.1-21.2. Contains a mouse E25 like gene, a Kinesin like pseudogene and ESTs /FEA=mRNA /DB_XREF=gi:2853186 /UG=Hs.17109 integral membrane protein 2A /FL=gb:AF038953.1 gb:NM_004867.1
216960_s_at	Consensus includes gb:AL049646 /DEF=Human DNA sequence from clone RP4-568F9 on chromosome 20 Contains the ZNF133 (zinc finger protein 133 (clone pHZ-13)) gene, part of a gene for a novel protein, ESTs, STSs, GSSs and CpG islands /FEA=mRNA_2 /DB_XREF=gi:11121205 /UG=Hs.78434 zinc finger protein 133 (clone pHZ-13)
213110_s_at	Consensus includes gb:AW052179 /FEA=EST /DB_XREF=gi:5914538 /DB_XREF=est:wx26h09.x1 /CLONE=IMAGE:2544833 /UG=Hs.169825 collagen, type IV, alpha 5 (Alport syndrome) /FL=gb:NM_000495.2
200918_s_at	gb:NM_003139.1 /DEF=Homo sapiens signal recognition particle receptor (docking protein) (SRPR), mRNA. /FEA=mRNA /GEN=SRPR /PROD=signal recognition particle receptor (dockingprotein) /DB_XREF=gi:4507222 /UG=Hs.75730 signal recognition particle receptor (docking protein) /FL=gb:BC001162.1 gb:NM_003139.1



Affy Probe ID	Sequence Description
209438_at	Consensus includes gb:AL096700 /DEF=Human DNA sequence from clone RP3-499B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serinethreonine protein phosphatase PPEF, the 3 end of the PHKA2 gene for liver phosphorylase kinase alpha 2, ESTs, STSs and GSSs /FEA=mRNA_2 /DB_XREF=gi:7077104 /UG=Hs.54941 phosphorylase kinase, alpha 2 (liver) /FL=gb:D38616.1 gb:NM_000292.1
200700_s_at	gb:NM_006854.2 /DEF=Homo sapiens KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 (KDEL2), mRNA. /FEA=mRNA /GEN=KDEL2 /PROD=KDEL receptor 2 /DB_XREF=gi:8051609 /UG=Hs.118778 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2 /FL=gb:NM_006854.2
202833_s_at	gb:NM_000295.1 /DEF=Homo sapiens serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 (SERPINA1), mRNA. /FEA=mRNA /GEN=SERPINA1 /PROD=serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /DB_XREF=gi:4505792 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1 /FL=gb:AF130068.1 gb:M11465.1 gb:K01396.1 gb:NM_000295.1
202607_at	Consensus includes gb:AL526632 /FEA=EST /DB_XREF=gi:12790125 /DB_XREF=est:AL526632 /CLONE=CS0DC020YC21 (3 prime) /UG=Hs.20894 N-deacetylaseN-sulfotransferase (heparan glucosaminyl) 1 /FL=gb:U36600.1 gb:NM_001543.1 gb:U17970.1 gb:U18918.1
221103_s_at	gb:NM_018338.1 /DEF=Homo sapiens hypothetical protein FLJ11142 (FLJ11142), mRNA. /FEA=mRNA. /GEN=FLJ11142 /PROD=hypothetical protein FLJ11142 /DB_XREF=gi:8922896 /UG=Hs.272244 hypothetical protein FLJ11142 /FL=gb:NM_018338.1
210544_s_at	gb:BC002430.1 /DEF=Homo sapiens, aldehyde dehydrogenase 10 (fatty aldehyde dehydrogenase), clone MGC:2136, mRNA, complete cds. /FEA=mRNA /PROD=aldehyde dehydrogenase 10 (fatty aldehydedehydrogenase) /DB_XREF=gi:12803234 /UG=Hs.159608 aldehyde dehydrogenase 3 family, member A2 /FL=gb:BC002430.1
221581_s_at	gb:AF257135.1 /DEF=Homo sapiens WBSCR15 protein (WBSCR15) mRNA, complete cds. /FEA=mRNA /GEN=WBSCR15 /PROD=WBSCR15 protein /DB_XREF=gi:9651998 /UG=Hs.56607 Williams-Beuren syndrome chromosome region 5 /FL=gb:NM_022040.1 gb:BC001609.1 gb:AF257135.1
219482_at	gb:NM_017438.1 /DEF=Homo sapiens chromosome 21 open reading frame 18 (C21ORF18), mRNA. /FEA=mRNA /GEN=C21ORF18 /PROD=chromosome 21 open reading frame 18 /DB_XREF=gi:8393012 /UG=Hs.50748 chromosome 21 open reading frame 18 /FL=gb:NM_017438.1

Affy Probe_ID	Sequence Description
218950_at	gb:NM_022481.1 /DEF=Homo sapiens hypothetical protein FLJ21065 (FLJ21065), mRNA. /FEA=mRNA /GEN=FLJ21065 /PROD=hypothetical protein FLJ21065 /DB_XREF=gi:11968032 /UG=Hs.25277 hypothetical protein FLJ21065 /FL=gb:NM_022481.1
212399_s_at	Consensus includes gb:D50911.2 /DEF=Homo sapiens mRNA for KIAA0121 protein, partial cds. /FEA=mRNA /GEN=KIAA0121 /PROD=KIAA0121 protein /DB_XREF=gi:6633996 /UG=Hs.155584 KIAA0121 gene product
219274_at	gb:NM_012338.1 /DEF=Homo sapiens transmembrane 4 superfamily member (tetraspan NET-2) (NET-2), mRNA. /FEA=mRNA /GEN=NET-2 /PROD=transmembrane 4 superfamily member (tetraspanNET-2) /DB_XREF=gi:6912527 /UG=Hs.16529 transmembrane 4 superfamily member (tetraspan NET-2) /FL=gb:AF124522.1 gb:NM_012338.1
202498_s_at	Consensus includes gb:BE550486 /FEA=EST /DB_XREF=gi:9792178 /DB_XREF=est:7a27c01.x1 /CLONE=IMAGE:3219936 /UG=Hs.7594 solute carrier family 2 (facilitated glucose transporter), member 3 /FL=gb:M20681.1 gb:NM_006931.1
221234_s_at	gb:NM_021813.1 /DEF=Homo sapiens BTB and CNC homology 1, basic leucine zipper transcription factor 2 (BACH2), mRNA. /FEA=mRNA /GEN=BACH2 /PROD=BTB and CNC homology 1, basic leucine zippertranscription factor 2 /DB_XREF=gi:13540489 /FL=gb:NM_021813.1
219968_at	gb:NM_016089.1 /DEF=Homo sapiens KRAB-zinc finger protein SZF1-1 (SZF1), mRNA. /FEA=mRNA /GEN=SZF1 /PROD=KRAB-zinc finger protein SZF1-1 /DB_XREF=gi:7706726 /UG=Hs.19585 KRAB-zinc finger protein SZF1-1 /FL=gb:AF114816.1 gb:NM_016089.1
201506_at	gb:NM_000358.1 /DEF=Homo sapiens transforming growth factor, beta-induced, 68kD (TGFB1), mRNA. /FEA=mRNA /GEN=TGFB1 /PROD=transforming growth factor, beta-induced, 68kD /DB_XREF=gi:4507466 /UG=Hs.118787 transforming growth factor, beta-induced, 68kD /FL=gb:BC000097.1 gb:BC004972.1 gb:M77349.1 gb:NM_000358.1
203420_at	gb:NM_016255.1 /DEF=Homo sapiens Autosomal Highly Conserved Protein (AHCP), mRNA. /FEA=mRNA /GEN=AHCP /PROD=Autosomal Highly Conserved Protein /DB_XREF=gi:7705267 /UG=Hs.95260 Autosomal Highly Conserved Protein /FL=gb:AF097027.1 gb:NM_016255.1
222171_s_at	Consensus includes gb:AK023792.1 /DEF=Homo sapiens cDNA FLJ13730 fis, clone PLACE3000136. /FEA=mRNA /DB_XREF=gi:10435834 /UG=Hs.10043 hypothetical protein FLJ13074

Affy Probe_ID	Sequence Description
221488_s_at	gb:AF230924.1 /DEF=Homo sapiens brain acetylcholinesterase putative membrane anchor mRNA, complete cds. /FEA=mRNA /PROD=brain acetylcholinesterase putative membraneanchor /DB_XREF=gi:7341254 /UG=Hs.107187
213178_s_at	divalent cation tolerant protein CUTA /FL=gb:AF230924.1 Consensus includes gb:AB028989.1 /DEF=Homo sapiens mRNA for KIAA1066 protein, partial cds. /FEA=mRNA /GEN=KIAA1066 /PROD=KIAA1066 protein /DB_XREF=gi:5689468 /UG=Hs.88500 mitogen-activated protein kinase 8 interacting protein 3
205239_at	gb:NM_001657.1 /DEF=Homo sapiens amphiregulin (schwannoma-derived growth factor) (AREG), mRNA. /FEA=mRNA /GEN=AREG /PROD=amphiregulin (schwannoma-derived growth factor) /DB_XREF=gi:4502198 /UG=Hs.270833 amphiregulin (schwannoma-derived growth factor) /FL=gb:M30704.1 gb:NM_001657.1 Consensus includes gb:BE379006 /FEA=EST /DB_XREF=gi:9324371 /DB_XREF=est:601236272F1
212463_at	/CLONE=IMAGE:3608680 /UG=Hs.99766 Homo sapiens mRNA; cDNA DKFZp564J0323 (from clone DKFZp564J0323)
202059_s_at	gb:NM_002264.1 /DEF=Homo sapiens karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA. /FEA=mRNA /GEN=KPNA1 /PROD=karyopherin alpha 1 /DB_XREF=gi:4504894 /UG=Hs.169149 karyopherin alpha 1 (importin alpha 5) /FL=gb:BC002374.1 gb:BC003009.1 gb:NM_002264.1 Consensus includes gb:AF124489.1 /DEF=Homo sapiens arfaptin-1b mRNA, alternatively spliced, complete cds. /FEA=CDS /PROD=arfaptin-1b /DB_XREF=gi:4761515 /UG=Hs.301064 arfaptin 1 /FL=gb:AF124489.1
214483_s_at	Consensus includes gb:N22903 /FEA=EST /DB_XREF=gi:1137053 /DB_XREF=est:yx66e04.sl
203327_at	/CLONE=IMAGE:266718 /UG=Hs.1508 insulin-degrading enzyme /FL=gb:M21188.1 gb:NM_004969.1 Consensus includes gb:AA749101 /FEA=EST /DB_XREF=gi:2789059 /DB_XREF=est:ny11d02.sl
214022_s_at	/CLONE=IMAGE:1271427 /UG=Hs.146360 interferon induced transmembrane protein 1 (9-27)
202912_at	gb:NM_001124.1 /DEF=Homo sapiens adrenomedullin (ADM), mRNA. /FEA=mRNA /GEN=ADM /PROD=adrenomedullin /DB_XREF=gi:4501944 /UG=Hs.394 adrenomedullin /FL=gb:NM_001124.1 gb:D14874.1
210385_s_at	gb:AF106037.1 /DEF=Homo sapiens adipocyte-derived leucine aminopeptidase mRNA, complete cds. /FEA=mRNA /PROD=adipocyte-derived leucine aminopeptidase /DB_XREF=gi:6381988 /UG=Hs.247043 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator /FL=gb:AF106037.1
203017_s_at	Consensus includes gb:R52678 /FEA=EST /DB_XREF=gi:814580 /DB_XREF=est:y99c07.sl /CLONE=IMAGE:41738 /UG=Hs.22587 KIAA0923 protein /FL=gb:AB023140.1 gb:NM_014021.1

Affy Probe_ID	Sequence Description
202668_at	Consensus includes gb:BF001670 /FEA=EST /DB_XREF=gi:10701945 /DB_XREF=est:7g91e06.x1 /CLONE=IMAGE:3313858 /UG=Hs.30942 ephrin-B2 /FL=gb:U81262.1 gb:Nm_004093.1 gb:U16797.1 gb:BC000069.1 /DEF=Homo sapiens, retinoic acid receptor responder (tazarotene induced) 2, clone MGC:1544, mRNA, complete cds. /FEA=mRNA /PROD=retinoic acid receptor responder (tazarotene induced) 2 /DB_XREF=gi:12652642 /UG=Hs.37682 retinoic acid receptor responder (tazarotene induced) 2 /FL=gb:BC000069.1 gb:U77594.1 gb:AB015632.1 gb:Nm_002889.2
204157_s_at	gb:Nm_025164.1 /DEF=Homo sapiens KIAA0999 protein (KIAA0999), mRNA. /FEA=mRNA /GEN=KIAA0999 /PROD=hypothetical protein FLJ12240 /DB_XREF=gi:13386467 /UG=Hs.4278 KIAA0999 protein /FL=gb:Nm_025164.1
200806_s_at	Consensus includes gb:BE256479 /FEA=EST /DB_XREF=gi:9126938 /DB_XREF=est:601111293F1 /CLONE=IMAGE:3352031 /UG=Hs.79037 heat shock 60kD protein 1 (chaperonin) /FL=gb:BC002676.1 gb:BC003030.1 gb:M34664.1 gb:M22382.1 gb:Nm_002156.1
213923_at	Consensus includes gb:AW005535 /FEA=EST /DB_XREF=gi:5854313 /DB_XREF=est:wz87d04.x1 /CLONE=IMAGE:2565799 /UG=Hs.155218 E1B-55kDa-associated protein 5
206522_at	gb:Nm_004668.1 /DEF=Homo sapiens maltase-glucoamylase (alpha-glucosidase) (MGAM), mRNA. /FEA=mRNA /GEN=MGAM /PROD=alpha-glucosidase /DB_XREF=gi:4758711 /UG=Hs.122785 maltase-glucoamylase (alpha-glucosidase) /FL=gb:AF016833.1 gb:Nm_004668.1
201041_s_at	gb:Nm_004417.2 /DEF=Homo sapiens dual specificity phosphatase 1 (DUSP1), mRNA. /FEA=mRNA /GEN=DUSP1 /PROD=dual specificity phosphatase 1 /DB_XREF=gi:7108342 /UG=Hs.171695 dual specificity phosphatase 1 /FL=gb:Nm_004417.2
208002_s_at	gb:Nm_007274.1 /DEF=Homo sapiens cytosolic acyl coenzyme A thioester hydrolase (HBACH), mRNA. /FEA=mRNA /GEN=HBACH /PROD=cytosolic acyl coenzyme A thioester hydrolase /DB_XREF=gi:6857796 /UG=Hs.8679 cytosolic acyl coenzyme A thioester hydrolase /FL=gb:U91316.1 gb:Nm_007274.1
200660_at	gb:Nm_005620.1 /DEF=Homo sapiens S100 calcium-binding protein A11 (calgizzarin) (S100A11), mRNA. /FEA=mRNA /GEN=S100A11 /PROD=S100 calcium-binding protein A11 /DB_XREF=gi:5032056 /UG=Hs.256290 S100 calcium-binding protein A11 (calgizzarin) /FL=gb:D49355.1 gb:BC001410.1 gb:D50374.1 gb:Nm_005620.1 gb:D38583.1

Affy Probe ID	Sequence Description
201532_at	gb:NM_002788.1 /DEF=Homo sapiens proteasome (prosome, macropain) subunit, alpha type, 3 (PSMA3), mRNA. /FEA=mRNA /GEN=PSMA3 /PROD=proteasome (prosome, macropain) subunit, alphatype, 3 /DB_XREF=gi:4506182 /UG=Hs.167106 proteasome (prosome, macropain) subunit, alpha type, 3 /FL=gb:BC005265.1 gb:NM_002788.1
208944_at	gb:D50683.1 /DEF=Homo sapiens mRNA for TGF-betaIIIR alpha, complete cds. /FEA=mRNA /PROD=TGF-betaIIIR alpha /DB_XREF=gi:1827474 /UG=Hs.82028 transforming growth factor, beta receptor II (70-80kD) /FL=gb:D50683.1 Consensus includes gb:AW150913 /FEA=EST /DB_XREF=gi:6198811 /DB_XREF=est:hg42a01.x1 /CLONE=IMAGE:2630184 /UG=Hs.90998 KIAA0128 protein; septin 2
212415_at	gb:NM_003151.1 /DEF=Homo sapiens signal transducer and activator of transcription 4 (STAT4), mRNA. /FEA=mRNA /GEN=STAT4 /PROD=signal transducer and activator of transcription4 /DB_XREF=gi:4507254 /UG=Hs.80642 signal transducer and activator of transcription 4 /FL=gb:L78440.1 gb:NM_003151.1
201543_s_at	gb:NM_020150.1 /DEF=Homo sapiens SAR1 protein (SAR1), mRNA. /FEA=mRNA /GEN=SAR1 /PROD=SAR1 protein /DB_XREF=gi:9910541 /UG=Hs.110796 SAR1 protein /FL=gb:AY008268.1 gb:AL136724.1 gb:AF261717.1 gb:NM_020150.1
212157_at	Consensus includes gb:BE048514 /FEA=EST /DB_XREF=gi:8365567 /DB_XREF=est:hr50a03.x1 /CLONE=IMAGE:3131884 /UG=Hs.1501 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
219558_at	gb:NM_024524.1 /DEF=Homo sapiens hypothetical protein FLJ20986 (FLJ20986), mRNA. /FEA=mRNA /GEN=FLJ20986 /PROD=hypothetical protein FLJ20986 /DB_XREF=gi:13375667 /UG=Hs.324507 hypothetical protein FLJ20986 /FL=gb:NM_024524.1
221603_at	gb:BC000467.1 /DEF=Homo sapiens, Similar to peroxisomal biogenesis factor 16, clone MGC:8550, mRNA, complete cds. /FEA=mRNA /PROD=Similar to peroxisomal biogenesis factor 16 /DB_XREF=gi:12653394 /UG=Hs.100915 peroxisomal biogenesis factor 16 /FL=gb:BC000467.1
212552_at	Consensus includes gb:BE617588 /FEA=EST /DB_XREF=gi:9888526 /DB_XREF=est:601441509T1 /CLONE=IMAGE:3845643 /UG=Hs.3618 hippocalcin-like 1
205945_at	gb:NM_000565.1 /DEF=Homo sapiens interleukin 6 receptor (IL6R), mRNA. /FEA=mRNA /GEN=IL6R /PROD=interleukin 6 receptor /DB_XREF=gi:4504672 /UG=Hs.193400 interleukin 6 receptor /FL=gb:NM_000565.1

**Affy Probe\_ID**

**Sequence Description**

209124_at	gb:U70451.1 /DEF=Human myeloid differentiation primary response protein MyD88 mRNA, complete cds. /FEA=mRNA /PROD=MyD88 /DB_XREF=gi:1763090 /UG=Hs.82116 myeloid differentiation primary response gene (88) /FL=gb:U70451.1 gb:U84408.1 gb:Nm_002468.1
205778_at	gb:Nm_005046.1 /DEF=Homo sapiens kallikrein 7 (chymotryptic, stratum corneum) (KLK7), mRNA. /FEA=mRNA /GEN=KLK7 /PROD=kallikrein 7 (chymotryptic, stratum corneum) /DB_XREF=gi:4826949 /UG=Hs.151254 kallikrein 7 (chymotryptic, stratum corneum) /FL=gb:Nm_005046.1 gb:L33404.1
200807_s_at	gb:Nm_002156.1 /DEF=Homo sapiens heat shock 60kD protein 1 (chaperonin) (HSPD1), mRNA. /FEA=mRNA /GEN=HSPD1 /PROD=heat shock 60kD protein 1 (chaperonin) /DB_XREF=gi:4504520 /UG=Hs.79037 heat shock 60kD protein 1 (chaperonin) /FL=gb:BC002676.1 gb:BC003030.1 gb:M34664.1 gb:M22382.1 gb:Nm_002156.1
219956_at	gb:Nm_007210.2 /DEF=Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetyl-galactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA. /FEA=mRNA /GEN=GALNT6 /PROD=polypeptide N-acetyl-galactosaminyltransferase 6 /DB_XREF=gi:13124893 /UG=Hs.151678 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetyl-galactosaminyltransferase 6 (GalNAc-T6) /FL=gb:Nm_007210.2
200803_s_at	gb:AF033095.1 /DEF=Homo sapiens testis enhanced gene transcript protein (TEGT) mRNA, complete cds. /FEA=mRNA /GEN=TEGT /PROD=testis enhanced gene transcript protein /DB_XREF=gi:2645728 /UG=Hs.74637 testis enhanced gene transcript (BAX inhibitor 1) /FL=gb:BC000916.1 gb:AF033095.1 gb:Nm_003217.1
219174_at	gb:Nm_025103.1 /DEF=Homo sapiens hypothetical protein FLJ22621 (FLJ22621), mRNA. /FEA=mRNA /GEN=FLJ22621 /PROD=hypothetical protein FLJ22621 /DB_XREF=gi:13376668 /UG=Hs.288617 hypothetical protein FLJ22621 /FL=gb:Nm_025103.1
218815_s_at	gb:Nm_018022.1 /DEF=Homo sapiens hypothetical protein FLJ10199 (FLJ10199), mRNA. /FEA=mRNA /GEN=FLJ10199 /PROD=hypothetical protein FLJ10199 /DB_XREF=gi:8922276 /UG=Hs.30925 hypothetical protein FLJ10199 /FL=gb:BC000202.1 gb:BC000593.1 gb:Nm_018022.1
209013_x_at	gb:AF091395.1 /DEF=Homo sapiens Trio isoform mRNA, complete cds. /FEA=mRNA /PROD=Trio isoform /DB_XREF=gi:3644047 /UG=Hs.171957 triple functional domain (PTPRF interacting) /FL=gb:AF091395.1
209933_s_at	gb:AF020314.1 /DEF=Homo sapiens CMRF-35-H9 mRNA, complete cds. /FEA=mRNA /PROD=CMRF-35-H9 /DB_XREF=gi:4103065 /UG=Hs.9688 leukocyte membrane antigen /FL=gb:AF020314.1
220126_at	gb:Nm_013270.1 /DEF=Homo sapiens testes-specific protease 50 (TSP50), mRNA. /FEA=mRNA /GEN=TSP50 /PROD=testes-specific protease 50 /DB_XREF=gi:7019562 /UG=Hs.120365 testes-specific protease 50 /FL=gb:AF100707.1 gb:Nm_013270.1

Affy Probe_ID	Sequence Description
221266_s_at	gb:NM_030788.1 /DEF=Homo sapiens DC-specific transmembrane protein (LOC81501), mRNA. /FEA=mRNA /GEN=LOC81501 /PROD=DC-specific transmembrane protein /DB_XREF=gi:13540564 /FL=gb:NM_030788.1
218194_at	gb:NM_015523.1 /DEF=Homo sapiens small fragment nuclease (DKFZP566E144), mRNA. /FEA=mRNA /GEN=DKFZP566E144 /PROD=small fragment nuclease /DB_XREF=gi:7661645 /UG=Hs.7527 small fragment nuclease /FL=gb:AF151872.1 gb:AL110239.1 gb:NM_015523.1
203860_at	gb:NM_000282.1 /DEF=Homo sapiens propionyl Coenzyme A carboxylase, alpha polypeptide (PCCA), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=PCCA /PROD=Propionyl-Coenzyme A carboxylase, alapholypeptide precursor /DB_XREF=gi:4557832 /UG=Hs.80741 propionyl Coenzyme A carboxylase, alpha polypeptide /FL=gb:BC000140.1 gb:NM_000282.1
205500_at	gb:NM_001735.1 /DEF=Homo sapiens complement component 5 (C5), mRNA. /FEA=mRNA /GEN=C5 /PROD=complement component 5 /DB_XREF=gi:4502506 /UG=Hs.1281 complement component 5 /FL=gb:M57729.1 gb:NM_001735.1
201192_s_at	gb:NM_006224.1 /DEF=Homo sapiens phosphatidylinositol transfer protein (PITPN), mRNA. /FEA=mRNA /GEN=PITPN /PROD=phosphatidylinositol transfer protein /DB_XREF=gi:5453907 /UG=Hs.79709 phosphatidylinositol transfer protein /FL=gb:D30036.1 gb:M73704.1 gb:NM_006224.1
217591_at	Consensus includes gb:BF725121 /FEA=EST /DB_XREF=gi:12041032 /DB_XREF=est:bx12e01.x1 /CLONE=bx12e01 /UG=Hs.272108 ESTs
203515_s_at	gb:NM_006556.1 /DEF=Homo sapiens phosphomevalonate kinase (PMVK), mRNA. /FEA=mRNA /GEN=PMVK /PROD=phosphomevalonate kinase /DB_XREF=gi:5729979 /UG=Hs.30954 phosphomevalonate kinase /FL=gb:L77213.1 gb:NM_006556.1
210976_s_at	gb:U24183.1 /DEF=Human phosphofructokinase (PFKM) mRNA, complete cds. /FEA=mRNA /GEN=PFKM /PROD=phosphofructokinase /DB_XREF=gi:975289 /UG=Hs.75160 phosphofructokinase, muscle /FL=gb:U24183.1
210115_at	gb:L05096.1 /DEF=Homo sapiens ribosomal protein L39 mRNA, complete cds. /FEA=mRNA /PROD=ribosomal protein L39 /DB_XREF=gi:388035 /UG=Hs.132748 Homo sapiens ribosomal protein L39 mRNA, complete cds /FL=gb:L05096.1
209653_at	gb:U93240.1 /DEF=Human importin alpha 3 mRNA, complete cds. /FEA=mRNA /PROD=importin alpha 3 /DB_XREF=gi:1928974 /UG=Hs.302499 karyopherin alpha 4 (importin alpha 3) /FL=gb:AB002533.2 gb:NM_002268.2 gb:U93240.1

**Affy Probe ID**

**Sequence Description**

208878_s_at	gb:AF092132.1 /DEF=Homo sapiens PAK2 mRNA, complete cds. /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds. /FL=gb:AF092132.1
216526_x_at	Consensus includes gb:AK024836.1 /DEF=Homo sapiens cDNA: FLJ21183 fis, clone CAS11634, highly similar to HSHLACW07 Homo sapiens mRNA for human leukocyte antigen C alpha chain. /FEA=mRNA
213556_at	/DB_XREF=gi:10437242 /UG=Hs.277477 major histocompatibility complex, class I, C
218966_at	Consensus includes gb:BE673445 /FEA=EST /DB_XREF=gi:10033986 /DB_XREF=est:7d35d10.x1 /CLONE=IMAGE:3249235 /UG=Hs.22049 Homo sapiens chromosome 19, cosmid R28379
205564_at	gb:NM_018728.1 /DEF=Homo sapiens myosin 5C (MYO5C), mRNA. /FEA=mRNA /GEN=MYO5C /PROD=myosin 5C /DB_XREF=gi:9055283 /UG=Hs.111782 myosin 5C /FL=gb:AF272390.1 gb:NM_018728.1
215646_s_at	gb:NM_007003.1 /DEF=Homo sapiens JM27 protein (JM27), mRNA. /FEA=mRNA /GEN=JM27 /PROD=JM27 protein /DB_XREF=gi:5901985 /UG=Hs.95420 JM27 protein /FL=gb:AJ005894.1 gb:NM_007003.1 gb:AF275258.1
219358_s_at	Consensus includes gb:R94644 /FEA=EST /DB_XREF=gi:970039 /DB_XREF=est:yq42a12.r1 /CLONE=IMAGE:198430 /UG=Hs.306542 Homo sapiens versican Vint isoform, mRNA, partial cds
219439_at	gb:NM_018404.1 /DEF=Homo sapiens centaurin-alpha 2 protein (HSA272195), mRNA. /FEA=mRNA /GEN=HSA272195 /PROD=centaurin-alpha 2 protein /DB_XREF=gi:8923762 /UG=Hs.28802 centaurin-alpha 2 protein /FL=gb:NM_018404.1
204818_at	gb:NM_020156.1 /DEF=Homo sapiens core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase (C1GALT1), mRNA. /FEA=mRNA /GEN=C1GALT1 /PROD=core1UDP-galactose:N-acetylgalactosamine-alpha-R beta1,3-galactosyltransferase /DB_XREF=gi:9910143 /UG=Hs.46744 core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase /FL=gb:AF155582.1 gb:NM_020156.1
207813_s_at	gb:NM_002153.1 /DEF=Homo sapiens hydroxysteroid (17-beta) dehydrogenase 2 (HSD17B2), mRNA. /FEA=mRNA /GEN=HSD17B2 /PROD=hydroxysteroid (17-beta) dehydrogenase 2 /DB_XREF=gi:4504502 /UG=Hs.155109 hydroxysteroid (17-beta) dehydrogenase 2 /FL=gb:L11708.1 gb:NM_002153.1
217971_at	gb:NM_004110.2 /DEF=Homo sapiens ferredoxin reductase (FDXR), transcript variant 2, nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=FDXR /PROD=ferredoxin reductase isoform 2 precursor /DB_XREF=gi:13435351 /UG=Hs.69745 ferredoxin reductase /FL=gb:NM_004110.2 gb:J03826.1
	gb:NM_021970.1 /DEF=Homo sapiens MEK partner 1 (MP1), mRNA. /FEA=mRNA /GEN=MP1 /PROD=MEK partner 1 /DB_XREF=gi:11496276 /UG=Hs.6361 MEK partner 1 /FL=gb:AF130115.1 gb:NM_021970.1 gb:AF201947.1



Affy Probe_ID	Sequence Description
210570_x_at	gb:U35002.1 /DEF=Human JNK2 beta1 protein kinase (JNK2B1) mRNA, complete cds. /FEA=mRNA /GEN=JNK2B1 /PROD=JNK2 beta1 protein kinase /DB_XREF=gi:1463132 /UG=Hs.246857 mitogen-activated protein kinase 9 /FL=gb:U35002.1 gb:U35003.1
201647_s_at	gb:NM_005506.1 /DEF=Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II) (CD36L2), mRNA. /FEA=mRNA /GEN=CD36L2 /PROD=CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II) /DB_XREF=gi:5031630 /UG=Hs.323567 CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II) /FL=gb:D12676.1 gb:NM_005506.1
218109_s_at	gb:NM_022736.1 /DEF=Homo sapiens hypothetical protein FLJ14153 (FLJ14153), mRNA. /FEA=mRNA /GEN=FLJ14153 /PROD=hypothetical protein FLJ14153 /DB_XREF=gi:12232392 /UG=Hs.7503 hypothetical protein FLJ14153 /FL=gb:NM_022736.1 gb:AB014732.1
205470_s_at	gb:NM_006853.1 /DEF=Homo sapiens kallikrein 11 (KLK11), mRNA. /FEA=mRNA /GEN=KLK11 /PROD=kallikrein 11 /DB_XREF=gi:5803198 /UG=Hs.57771 kallikrein 11 /FL=gb:AB012917.1 gb:NM_006853.1 gb:AB041036.1
208893_s_at	gb:BC005047.1 /DEF=Homo sapiens, clone MGC:12852, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12852) /DB_XREF=gi:13477170 /UG=Hs.180383 dual specificity phosphatase 6 /FL=gb:BC003562.1 gb:BC003143.1 gb:BC005047.1 gb:AB013382.1 gb:NM_001946.1
212771_at	Consensus includes gb:AU150943 /FEA=EST /DB_XREF=gi:11012464 /DB_XREF=est:AU150943 /CLONE=NT2RP2003984 /UG=Hs.66762 Homo sapiens mRNA; cDNA DKFZp564A026 (from clone DKFZp564A026)
203821_at	gb:NM_001945.1 /DEF=Homo sapiens diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor) (DTR), mRNA. /FEA=mRNA /GEN=DTR /PROD=diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor) /DB_XREF=gi:4503412 /UG=Hs.799 diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor) /FL=gb:M60278.1 gb:NM_001945.1
202620_s_at	gb:NM_000935.1 /DEF=Homo sapiens procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2 (PLOD2), mRNA. /FEA=mRNA /GEN=PLOD2 /PROD=procollagen-lysine, 2-oxoglutarate 5-dioxygenase(lysine hydroxylase) 2 /DB_XREF=gi:4505888 /UG=Hs.41270 procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2 /FL=gb:U84573.1 gb:NM_000935.1

**Affy Probe ID**

**Sequence Description**

203973_s_at	gb:NM_005195.1 /DEF=Homo sapiens CCAATenhancer binding protein (CEBP), delta (CEBPD), mRNA. /FEA=mRNA /GEN=CEBPD /PROD=CCAATenhancer binding protein (CEBP), delta /DB_XREF=gi:4885130 /UG=Hs.76722 CCAATenhancer binding protein (CEBP), delta /FL=gb:M83667.1 gb:NM_005195.1 Consensus includes gb:BF674349 /FEA=EST /DB_XREF=gi:11948244 /DB_XREF=est:602137094F1 /CLONE=IMAGE:4273679 /UG=Hs.24341-transcriptional co-activator with PDZ-binding motif (TAZ) /FL=gb:NM_015472.1
202133_at	Consensus includes gb:AF007150.1 /DEF=Homo sapiens clone 23767 and 23782 mRNA sequences. /FEA=mRNA /DB_XREF=gi:2852628 /UG=Hs.8025 Homo sapiens clone 23767 and 23782 mRNA sequences
211564_s_at	gb:BC003096.1 /DEF=Homo sapiens, Similar to LIM domain protein, clone MGC:1645, mRNA, complete cds. /FEA=mRNA /PROD=Similar to LIM domain protein /DB_XREF=gi:13111856 /UG=Hs.79691 LIM domain protein /FL=gb:BC003096.1
210122_at	gb:BC005303.1 /DEF=Homo sapiens, Similar to protamine 2, clone MGC:12373, mRNA, complete cds. /FEA=mRNA /PROD=Similar to protamine 2 /DB_XREF=gi:13529034 /UG=Hs.2324 protamine 2 /FL=gb:BC005303.1 gb:NM_002762.1
222369_at	Consensus includes gb:AW971254 /FEA=EST /DB_XREF=gi:8161099 /DB_XREF=est:EST383343 /UG=Hs.178433 ESTs
214651_s_at	Consensus includes gb:U41813.1 /DEF=Human class I homeoprotein (HOXA9) mRNA, partial cds: /FEA=mRNA /GEN=HOXA9 /PROD=HOXA9 /DB_XREF=gi:1184168 /UG=Hs.127428 homeo box A9 /FL=gb:NM_002142.1
211528_x_at	gb:M90685.1 /DEF=Human lymphocyte antigen (HLA-G2.2) mRNA, complete cds. /FEA=mRNA /GEN=HLA-G2.2 /PROD=b2 microglobulin /DB_XREF=gi:184211 /UG=Hs.73885 HLA-G histocompatibility antigen, class I, G /FL=gb:M90685.1
201431_s_at	gb:NM_001387.1 /DEF=Homo sapiens dihydropyrimidinase-like 3 (DPYSL3), mRNA. /FEA=mRNA /GEN=DPYSL3 /PROD=dihydropyrimidinase-like 3 /DB_XREF=gi:4503378 /UG=Hs.74566 dihydropyrimidinase-like 3 /FL=gb:D78014.1 gb:NM_001387.1
222062_at	Consensus includes gb:AI983115 /FEA=EST /DB_XREF=gi:5810334 /DB_XREF=est:wu18b02.x1 /CLONE=IMAGE:2517291 /UG=Hs.132781 class I cytokine receptor
213479_at	Consensus includes gb:U26662.1 /DEF=Human neuronal pentraxin II (NPTX2) mRNA, partial cds. /FEA=mRNA /GEN=NPTX2 /PROD=neuronal pentraxin II /DB_XREF=gi:881933 /UG=Hs.3281 neuronal pentraxin II

Affy Probe_ID	Sequence Description
219683_at	gb:NM_017412.1 /DEF=Homo sapiens frizzled (Drosophila) homolog 3 (FZD3), mRNA. /FEA=mRNA /GEN=FZD3 /PROD=frizzled (Drosophila) homolog 3 /DB_XREF=gi:8393377 /UG=Hs.40735 frizzled (Drosophila) homolog 3 /FL=gb:AY005130.3 gb:AB039723.1 gb:NM_017412.1
202890_at	Consensus includes gb:AW242297 /FEA=EST /DB_XREF=gi:6576051 /DB_XREF=est:xm96b11.x1 /CLONE=IMAGE:2692029 /UG=Hs.146388 microtubule-associated protein 7 /FL=gb:NM_003980.1
201147_s_at	Consensus includes gb:BF347089 /FEA=EST /DB_XREF=gi:11294684 /DB_XREF=est:602020842F1 /CLONE=IMAGE:4156354 /UG=Hs.245188 tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory) /FL=gb:U67195.1 gb:U02571.1 gb:U14394.1 gb:NM_000362.2
214721_x_at	Consensus includes gb:AL162074.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762L106 (from clone DKFZp762L106); partial cds. /FEA=mRNA /GEN=DKFZp762L106 /PROD=hypothetical protein /DB_XREF=gi:7328153 /UG=Hs.3903 Cdc42 effector protein 4; binder of Rho GTPases 4
200968_s_at	gb:NM_000942.1 /DEF=Homo sapiens peptidylprolyl isomerase B (cyclophilin B) (PPIB), mRNA. /FEA=mRNA /GEN=PPIB /PROD=peptidylprolyl isomerase B (cyclophilin B) /DB_XREF=gi:4758949 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin B) /FL=gb:BC001125.1 gb:M60857.1 gb:M63573.1 gb:NM_000942.1
212430_at	Consensus includes gb:AL109955 /DEF=Human DNA sequence from clone RP4-800J21 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 3 part of the RAE1 gene for a homolog to RNA export protein 1 from S.pombe and the gene for the ssDNA binding protein SEB4D.n /FEA=mRNA_1 /DB_XREF=gi:11558768 /UG=Hs.236361 seb4D
59644_at	Cluster Incl. AI735391:at10e09.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2354728 /clone_end=3 /gb=AI735391 /gi=5056915 /ug=Hs.20137 /len=567
214912_at	Consensus includes gb:AK022067.1 /DEF=Homo sapiens cDNA FLJ12005 fis, clone HEMBB1001565. /FEA=mRNA /DB_XREF=gi:10433387 /UG=Hs.287477 Homo sapiens cDNA FLJ12005 fis, clone HEMBB1001565
203894_at	gb:NM_016437.1 /DEF=Homo sapiens tubulin, gamma 2 (TUBG2), mRNA. /FEA=mRNA /GEN=TUBG2 /PROD=tubulin, gamma 2 /DB_XREF=gi:7706750 /UG=Hs.279669 tubulin, gamma 2 /FL=gb:AF225971.1 gb:NM_016437.1
211530_x_at	gb:M90686.1 /DEF=Human lymphocyte antigen (HLA-G3) mRNA, complete cds. /FEA=mRNA /GEN=HLA-G3 /PROD=b2 microglobulin /DB_XREF=gi:184213 /UG=Hs.73885 HLA-G histocompatibility antigen, class I, G /FL=gb:M90686.1

**Affy Probe ID**

**Sequence Description**

222056_s_at	Consensus includes gb:AA723370 /FEA=EST /DB_XREF=gi:2741077 /DB_XREF=est:zg73a01.s1 /CLONE=IMAGE:398952 /UG=Hs.279932 CGI-105 protein
209099_x_at	gb:U73936.1 /DEF=Homo sapiens Jagged 1 (HJ1) mRNA, complete cds. /FEA=mRNA /GEN=HJ1 /PROD=Jagged 1 /DB_XREF=gi:1695273 /UG=Hs.91143 jagged 1 (Alagille syndrome) /FL=gb:U61276.1 gb:U73936.1 gb:AF003837.1 gb:AF028593.1 gb:NM_000214.1
218193_s_at	gb:NM_016072.1 /DEF=Homo sapiens CGI-141 protein (LOC51026), mRNA. /FEA=mRNA /GEN=LOC51026 /PROD=CGI-141 protein /DB_XREF=gi:7705635 /UG=Hs.62275 CGI-141 protein /FL=gb:AF151899.1 gb:AL136571.1 gb:NM_016072.1
205066_s_at	gb:NM_006208.1 /DEF=Homo sapiens ectonucleotide pyrophosphatasephosphodiesterase 1 (ENPPI), mRNA. /FEA=mRNA /GEN=ENPPI /PROD=ectonucleotide pyrophosphatasephosphodiesterase 1 /DB_XREF=gi:13324676 /UG=Hs.11951 ectonucleotide pyrophosphatasephosphodiesterase 1 /FL=gb:NM_006208.1 gb:M57736.1
209710_at	Consensus includes gb:AL563460 /FEA=EST /DB_XREF=gi:12912874 /DB_XREF=est:AL563460 /CLONE=CS0DD006YG06 (3 prime) /UG=Hs.760 GATA-binding protein 2 /FL=gb:M68891.1
220402_at	gb:NM_022112.1 /DEF=Homo sapiens p53-regulated apoptosis-inducing protein 1 (P53AIP1), mRNA. /FEA=mRNA /GEN=P53AIP1 /PROD=p53-regulated apoptosis-inducing protein 1 /DB_XREF=gi:11545826 /UG=Hs.160953 p53-regulated apoptosis-inducing protein 1 /FL=gb:AB045832.1 gb:NM_022112.1
209295_at	gb:AF016266.1 /DEF=Homo sapiens TRAIL receptor 2 mRNA, complete cds. /FEA=mRNA /PROD=TRAIL receptor 2 /DB_XREF=gi:2529562 /UG=Hs.51233 tumor necrosis factor receptor superfamily, member 10b /FL=gb:BC001281.1 gb:AF018658.1 gb:AF016849.1 gb:AF016266.1
203395_s_at	gb:NM_005524.2 /DEF=Homo sapiens hairy (Drosophila)-homolog (HRY), mRNA. /FEA=mRNA /GEN=HRY /PROD=hairy (Drosophila)-homolog /DB_XREF=gi:8400709 /UG=Hs.250666 hairy (Drosophila)-homolog /FL=gb:AF264785.1 gb:NM_005524.2
222116_s_at	Consensus includes gb:AL157485.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762O207 (from clone DKFZp762O207). /FEA=mRNA /DB_XREF=gi:7018528 /UG=Hs.91973 hypothetical protein
202723_s_at	Consensus includes gb:AW117498 /FEA=EST /DB_XREF=gi:6086082 /DB_XREF=est:xd92e10.x1 /CLONE=IMAGE:2605098 /UG=Hs.170133 forkhead box O1A (rhabdomyosarcoma) /FL=gb:AF032885.1 gb:U02310.1 gb:NM_002015.2

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203687\_at

**Sequence Description**

gb:NM\_002996.1 /DEF=Homo sapiens small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin) (SCYD1), mRNA. /FEA=mRNA /GEN=SCYD1 /PROD=small inducible cytokine subfamily D(Cys-X3-Cys), member 1 (fractalkine, neurotactin) /DB\_XREF=gi:4506856 /UG=Hs.80420 small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin) /FL=gb:BC001163.1 gb:U91835.1 gb:NM\_002996.1 gb:NM\_019598.1 /DEF=Homo sapiens kallikrein 12 (KLK12), mRNA. /FEA=mRNA /GEN=KLK12 /PROD=kallikrein 12 /DB\_XREF=gi:9665235 /UG=Hs.159679 kallikrein 12 /FL=gb:NM\_019598.1 gb:NM\_000130.2 /DEF=Homo sapiens coagulation factor V (proaccelerin, labile factor) (F5), mRNA. /FEA=mRNA /GEN=F5 /PROD=coagulation factor V precursor /DB\_XREF=gi:10518500 /UG=Hs.30054 coagulation factor V (proaccelerin, labile factor) /FL=gb:NM\_000130.2 gb:M16967.1 gb:M14335.1

Consensus includes gb:NM\_006713.1 /DEF=Homo sapiens activated RNA polymerase II transcription cofactor 4 (PC4), mRNA. /FEA=CDS /GEN=PC4 /PROD=activated RNA polymerase II transcriptioncofactor 4 /DB\_XREF=gi:5729967 /UG=Hs.74861 activated RNA polymerase II transcription cofactor 4 /FL=gb:U12979.1 gb:NM\_006713.1 gb:AF288571.1 /DEF=Homo sapiens lymphoid enhancer factor-1 (LEF1) mRNA, complete cds. /FEA=mRNA /GEN=LEF1 /PROD=lymphoid enhancer factor-1 /DB\_XREF=gi:9858157 /UG=Hs.44865 lymphoid enhancer binding factor-1 /FL=gb:AF198532.1 gb:NM\_016269.1 gb:AF288571.1 gb:NM\_005196.1 /DEF=Homo sapiens centromere protein F (350400kD, mitotin) (CENPF), mRNA. /FEA=mRNA /GEN=CENPF /PROD=centromere protein F (350400kD, mitotin) /DB\_XREF=gi:4885132 /UG=Hs.77204 centromere protein F (350400kD, mitotin) /FL=gb:NM\_005196.1 gb:U19769.1

Consensus includes gb:BC004912.1 /DEF=Homo sapiens, clone IMAGE:3534745, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3534745) /DB\_XREF=gi:13436208 /UG=Hs.198689 KIAA0728 protein

Consensus includes gb:BF110411 /FEA=EST /DB\_XREF=gi:10940101 /DB\_XREF=est:7n52f02.x1 /CLONE=IMAGE:3568274 /UG=Hs.16959 ESTs

gb:NM\_022168.1 /DEF=Homo sapiens melanoma differentiation associated protein-5 (MDA5), mRNA. /FEA=mRNA /GEN=MDA5 /PROD=melanoma differentiation associated protein-5 /DB\_XREF=gi:11545921 /UG=Hs.293591 melanoma differentiation associated protein-5 /FL=gb:AF095844.1 gb:NM\_022168.1 gb:AY017378.1

Affy Probe ID	Sequence Description
219489_s_at	gb:NM_017821.1 /DEF=Homo sapiens hypothetical protein FLJ20435 (FLJ20435), mRNA. /FEA=mRNA /GEN=FLJ20435 /PROD=hypothetical protein FLJ20435 /DB_XREF=gi:8923408 /UG=Hs.11408 hypothetical protein FLJ20435 /FL=gb:NM_017821.1
207352_s_at	gb:NM_021911.1 /DEF=Homo sapiens gamma-aminobutyric acid (GABA) A receptor, beta 2 (GABRB2), transcript variant 1, mRNA. /FEA=mRNA /GEN=GABRB2 /PROD=gamma-aminobutyric acid (GABA) A receptor, beta2, isoform 1 /DB_XREF=gi:12548784 /UG=Hs.103998 gamma-aminobutyric acid (GABA) A receptor, beta 2 /FL=gb:NM_021911.1
206605_at	gb:NM_006025.1 /DEF=Homo sapiens protease, serine, 22 (P11), mRNA. /FEA=mRNA /GEN=P11 /PROD=protease, serine, 22 /DB_XREF=gi:5174622 /UG=Hs.997 protease, serine, 22 /FL=gb:M32402.1 gb:NM_006025.1
203753_at	gb:NM_003199.1 /DEF=Homo sapiens transcription factor 4 (TCF4), mRNA. /FEA=mRNA /GEN=TCF4 /PROD=transcription factor 4, isoform b /DB_XREF=gi:4507398 /UG=Hs.326198 transcription factor 4 /FL=gb:M74719.1 gb:NM_003199.1
217999_s_at	Consensus includes gb:NM_007350.1 /DEF=Homo sapiens pleckstrin homology-like domain, family A, member 1 (PHLDA1), mRNA. /FEA=mRNA /GEN=PHLDA1 /PROD=pleckstrin homology-like domain, family A, member 1 /DB_XREF=gi:6679302 /UG=Hs.82101 pleckstrin homology-like domain, family A, member 1 /FL=gb:NM_007350.1
208188_at	gb:NM_000226.1 /DEF=Homo sapiens keratin 9 (epidermolytic palmoplantar keratoderma) (KRT9), mRNA. /FEA=mRNA /GEN=KRT9 /PROD=keratin 9 /DB_XREF=gi:4557704 /UG=Hs.2783 keratin 9 (epidermolytic palmoplantar keratoderma) /FL=gb:NM_000226.1
211685_s_at	gb:AF251061.1 /DEF=Homo sapiens neurocalcin mRNA, complete cds. /FEA=mRNA /PROD=neurocalcin /DB_XREF=gi:13625183 /FL=gb:AF251061.1
209691_s_at	gb:BC003541.1 /DEF=Homo sapiens, hypothetical protein FLJ10488, clone MGC:10806, mRNA, complete cds. /FEA=mRNA /PROD=hypothetical protein FLJ10488 /DB_XREF=gi:13097653 /UG=Hs.279832 hypothetical protein FLJ10488 /FL=gb:BC003541.1
213362_at	Consensus includes gb:N73931 /FEA=EST /DB_XREF=gi:1231216 /DB_XREF=est:za74a07.s1 /CLONE=IMAGE:298260 /UG=Hs.323079 Homo sapiens mRNA; cDNA DKFZp564P116 (from clone DKFZp564P116)

**Affy Probe\_ID**

**Sequence Description**

203179_at	gb:NM_000155.1 /DEF=Homo sapiens galactose-1-phosphate uridylyltransferase (GALT), mRNA. /FEA=mRNA /GEN=GALT /PROD=galactose-1-phosphate uridylyltransferase /DB_XREF=gi:4557614 /UG=Hs.75641 galactose-1-phosphate uridylyltransferase /FL=gb:M60091.1 gb:NM_000155.1
208812_x_at	gb:BC004489.1 /DEF=Homo sapiens, major histocompatibility complex, class I, C, clone MGC:11039, mRNA, complete cds. /FEA=mRNA /PROD=major histocompatibility complex, class I, C /DB_XREF=gi:13325360 /UG=Hs.277477 major histocompatibility complex, class I, C /FL=gb:NM_002117.1 gb:M99388.1 gb:U06487.1 gb:BC002463.1 gb:BC004489.1 gb:D64145.1 gb:D38526.1 gb:D49552.1 gb:D49819.1 gb:M24097.1 gb:M84171.1 gb:M84172.1 gb:M84173.1 gb:M84174.1 gb:M26429.1 gb:M26430.1 gb:M26431.1 gb:U41420.1 gb:U41386.1 gb:D50852.1 gb:D50853.1 gb:D50854.1 gb:D83031.1 gb:U57028.1 gb:U06695.1 gb:U06696.1 gb:M99389.1 gb:M99390.1 gb:M28160.1 gb:U09853.1 gb:AF168611.1 gb:L38251.1 gb:D31817.1 Consensus includes gb:AU147799 /FEA=EST /DB_XREF=gi:11009320 /DB_XREF=est:AU147799 /CLONE=MAMMA1001744 /UG=Hs.288467 Homo sapiens cDNA FLJ12280 fis, clone MAMMA1001744 Consensus includes gb:AK025862.1 /DEF=Homo sapiens cDNA: FLJ22209 fis, clone HRC01496. /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA: FLJ22209 fis, clone HRC01496
213909_at	gb:NM_001778.1 /DEF=Homo sapiens CD48 antigen (B-cell membrane protein) (CD48), mRNA. /FEA=mRNA /GEN=CD48 /PROD=CD48 antigen (B-cell membrane protein) /DB_XREF=gi:4502674 /UG=Hs.901 CD48 antigen (B-cell membrane protein) /FL=gb:M59904.1 gb:M37766.1 gb:NM_001778.1
216449_x_at	gb:NM_001446.1 /DEF=Homo sapiens fatty acid binding protein 7, brain (FABP7), mRNA. /FEA=mRNA /GEN=FABP7 /PROD=fatty acid binding protein 7, brain /DB_XREF=gi:4557584 /UG=Hs.26770 fatty acid binding protein 7, brain /FL=gb:D50373.1 gb:U51338.1 gb:D88648.1 gb:U81235.1 gb:NM_001446.1
204118_at	Consensus includes gb:AK026980.1 /DEF=Homo sapiens cDNA: FLJ23327 fis, clone HEP12630, highly similar to HSZNF37 Homo sapiens ZNF37A mRNA for zinc finger protein. /FEA=mRNA /DB_XREF=gi:10439974 /UG=Hs.278064 Homo sapiens cDNA: FLJ23327 fis, clone HEP12630, highly similar to HSZNF37 Homo sapiens ZNF37A mRNA for zinc finger protein
205030_at	gb:NM_020202.1 /DEF=Homo sapiens Nit protein 2 (NIT2), mRNA. /FEA=mRNA /GEN=NIT2 /PROD=Nit protein 2 /DB_XREF=gi:9910459 /UG=Hs.15627 Nit protein 2 /FL=gb:AF260334.1 gb:AF284574.1 gb:NM_020202.1
215358_x_at	
218557_at	

**Affy Probe ID**

**Sequence Description**

203124_s_at	gb:NM_000617.1 /DEF=Homo sapiens solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 (SLC11A2), mRNA. /FEA=mRNA /GEN=SLC11A2 /PROD=solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 /DB_XREF=gi:10835168 /UG=Hs.57435 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 /FL=gb:NM_000617.1 gb:BC002592.1 gb:AB004857.1 gb:AF046997.1
219305_x_at	gb:NM_012168.1 /DEF=Homo sapiens F-box only protein 2 (FBXO2), mRNA. /FEA=mRNA /GEN=FBXO2 /PROD=F-box only protein 2 /DB_XREF=gi:6912363 /UG=Hs.132753 F-box only protein 2 /FL=gb:NM_012168.1
205807_s_at	gb:NM_020127.1 /DEF=Homo sapiens tuftelin 1 (TUFT1), mRNA. /FEA=mRNA /GEN=TUFT1 /PROD=tuftelin 1 /DB_XREF=gi:9910595 /UG=Hs.283009 tuftelin 1 /FL=gb:AL136917.1 gb:AF254260.1 gb:NM_020127.1
42361_g_at	Cluster Incl. AI588986:tkl15g03.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2151124 /clone_end=3 /gb=AI588986 /gi=4598034 /ug=Hs.236209 /len=499
213224_s_at	Consensus includes gb:AK025724.1 /DEF=Homo sapiens cDNA: FLJ22071 fis, clone HEP11691. /FEA=mRNA /DB_XREF=gi:10438333 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, clone HEP11691
210142_x_at	gb:AF117234.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB_XREF=gi:6563241 /UG=Hs.179986 flotillin 1 /FL=gb:AF117234.1
214297_at	Consensus includes gb:BE857703 /FEA=EST /DB_XREF=gi:10371993 /DB_XREF=est:7g46a02.x1 /CLONE=IMAGE:3309482 /UG=Hs.9004 chondroitin sulfate proteoglycan 4 (melanoma-associated)
203555_at	gb:NM_014369.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 18 (brain-derived) (PTPN18), mRNA. /FEA=mRNA /GEN=PTPN18 /PROD=protein tyrosine phosphatase, non-receptor type18 (brain-derived) /DB_XREF=gi:7657483 /UG=Hs.278597 protein tyrosine phosphatase, non-receptor type 18 (brain-derived) /FL=gb:NM_014369.1
211464_x_at	gb:U20537.1 /DEF=Human cysteine protease Mch2 isoform beta (Mch2) mRNA, complete cds. /FEA=mRNA /GEN=Mch2 /PROD=cysteine protease Mch2 isoform beta /DB_XREF=gi:882255 /UG=Hs.3280 caspase 6, apoptosis-related cysteine protease /FL=gb:U20537.1
221521_s_at	gb:BC003186.1 /DEF=Homo sapiens, HSPC037 protein, clone MGC:673, mRNA, complete cds. /FEA=mRNA /PROD=HSPC037 protein /DB_XREF=gi:13112024 /UG=Hs.108196 HSPC037 protein /FL=gb:BC003186.1 gb:AF201939.1
207146_at	gb:NM_002278.1 /DEF=Homo sapiens keratin, hair, acidic,2 (KRTHA2), mRNA. /FEA=mRNA /GEN=KRTHA2 /PROD=type I hair keratin 2 /DB_XREF=gi:4504922 /UG=Hs.41752 keratin, hair, acidic,2 /FL=gb:NM_002278.1



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**Sequence Description**

219880\_at gb:NM\_022907.1 /DEF=Homo sapiens hypothetical protein FLJ23053 (FLJ23053), mRNA. /FEA=mRNA /GEN=FLJ23053 /PROD=hypothetical protein FLJ23053 /DB\_XREF=gi:12597650 /UG=Hs.94037 hypothetical protein FLJ23053 /FL=gb:NM\_022907.1

212298\_at Consensus includes gb:BE620457 /FEA=EST /DB\_XREF=gi:9891395 /DB\_XREF=est:601483690F1 /CLONE=IMAGE:3886055 /UG=Hs.69285 neuropilin 1 /FL=gb:AF018956.1 gb:AF016050.1 gb:NM\_003873.1

212990\_at Consensus includes gb:AB020717.1 /DEF=Homo sapiens mRNA for KIAA0910 protein, partial cds. /FEA=mRNA /GEN=KIAA0910 /PROD=KIAA0910 protein /DB\_XREF=gi:4240308 /UG=Hs.127416 synaptotagmin 1

210305\_at gb:AB042557.1 /DEF=Homo sapiens mRNA, similar to rat myomegalin, complete cds. /FEA=mRNA /DB\_XREF=gi:11138039 /UG=Hs.265848 similar to rat myomegalin /FL=gb:AB042557.1

217311\_at Consensus includes gb:AF086790.1 /DEF=Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEN=ACON /PROD=aconitase precursor /DB\_XREF=gi:3600097 /UG=Hs.166002 Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds

209539\_at Consensus includes gb:D25304.1 /DEF=Human mRNA for KIAA0006 gene, partial cds. /FEA=mRNA /GEN=KIAA0006 /DB\_XREF=gi:435445 /UG=Hs.79307 RacCdc42 guanine exchange factor (GEF) 6 /FL=gb:D13631.1

200776\_s\_at Consensus includes gb:AL518328 /FEA=EST /DB\_XREF=gi:12781821 /DB\_XREF=est:AL518328 /CLONE=CS0DA009YK18 (3 prime) /UG=Hs.155291 KIAA0005 gene product /FL=gb:D13630.1 gb:NM\_014670.1

217699\_at Consensus includes gb:AV700338 /FEA=EST /DB\_XREF=gi:10302309 /DB\_XREF=est:AV700338 /CLONE=GKCAMH12 /UG=Hs.286243 ESTs

202716\_at gb:NM\_002827.1 /DEF=Homo sapiens protein tyrosine phosphatase, non-receptor type 1 (PTPN1), mRNA. /FEA=mRNA /GEN=PTPN1 /PROD=protein tyrosine phosphatase, non-receptor type1 /DB\_XREF=gi:4506288 /UG=Hs.155894 protein tyrosine phosphatase, non-receptor type 1 /FL=gb:M33689.1 gb:M31724.1 gb:NM\_002827.1

211787\_s\_at gb:BC006210.1 /DEF=Homo sapiens, Similar to eukaryotic translation initiation factor 4A, isoform 1, clone MGC:8484, mRNA, complete cds. /FEA=mRNA /PROD=Similar to eukaryotic translation initiation factor 4A, isoform 1 /DB\_XREF=gi:13623224 /FL=gb:BC006210.1

215985\_at Consensus includes gb:X92110.1 /DEF=H.sapiens mRNA for hcgVIII protein. /FEA=mRNA /DB\_XREF=gi:1216163 /UG=Hs.153618 HCGVIII-1 protein

**Affy Probe ID**

**Sequence Description**

216442\_x\_at Consensus includes gb:AK026737.1 /DEF=Homo sapiens cDNA: FLJ23084 fis, clone LNG06602, highly similar to HSF1B1 Human mRNA for fibronectin (FN precursor). /FEA=mRNA /DB\_XREF=gi:10439658 /UG=Hs.287820 fibronectin 1

206111\_at gb:NM\_002934.1 /DEF=Homo sapiens ribonuclease, RNase A family, 2 (liver, eosinophil-derived neurotoxin) (RNASE2), mRNA. /FEA=mRNA /GEN=RNASE2 /PROD=ribonuclease, RNase A family, 2 (liver, eosinophil-derived neurotoxin) /DB\_XREF=gi:4506548 /UG=Hs.728 ribonuclease, RNase A family, 2 (liver, eosinophil-derived neurotoxin) /FL=gb:M24157.1 gb:M30510.1 gb:NM\_002934.1 gb:M28129.1

208044\_s\_at gb:NM\_006238.1 /DEF=Homo sapiens peroxisome proliferative activated receptor, delta (PPARD), mRNA. /FEA=mRNA /GEN=PPARD /PROD=peroxisome proliferative activated receptor, delta /DB\_XREF=gi:5453939 /UG=Hs.106415 peroxisome proliferative activated receptor, delta /FL=gb:L07592.1 gb:NM\_006238.1

202579\_x\_at gb:NM\_006353.1 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein 17-like 3 (HMG17L3), mRNA. /FEA=mRNA /GEN=HMG17L3 /PROD=high-mobility group (nonhistone chromosomal) protein 17-like 3 /DB\_XREF=gi:10835239 /UG=Hs.236774 high-mobility group (nonhistone chromosomal) protein 17-like 3 /FL=gb:NM\_006353.1 gb:U90549.1

218853\_s\_at gb:NM\_019556.1 /DEF=Homo sapiens hypothetical protein dJ473B4 (DJ473B4), mRNA. /FEA=mRNA /GEN=DJ473B4 /PROD=hypothetical protein dJ473B4 /DB\_XREF=gi:9506542 /UG=Hs.57549 hypothetical protein dJ473B4 /FL=gb:NM\_019556.1

218245\_at gb:NM\_015516.1 /DEF=Homo sapiens hypothetical protein, estradiol-induced (E2IG4), mRNA. /FEA=mRNA /GEN=E2IG4 /PROD=hypothetical protein, estradiol-induced /DB\_XREF=gi:7661667 /UG=Hs.8361 hypothetical protein, estradiol-induced /FL=gb:AF191019.1 gb:NM\_015516.1

202974\_at gb:NM\_002436.2 /DEF=Homo sapiens membrane protein, palmitoylated 1 (55kD) (MPP1), mRNA. /FEA=mRNA /GEN=MPP1 /PROD=palmitoylated membrane protein 1 /DB\_XREF=gi:6006024 /UG=Hs.1861 membrane protein, palmitoylated 1 (55kD) /FL=gb:BC002392.1 gb:M64925.1 gb:NM\_002436.2

212127\_at Consensus includes gb:BE379408 /FEA=EST /DB\_XREF=gi:9324773 /DB\_XREF=est:601237213F1 /CLONE=IMAGE:3609315 /UG=Hs.183800 Ran GTPase activating protein 1 /FL=gb:NM\_002883.1

201798\_s\_at gb:NM\_013451.1 /DEF=Homo sapiens fer-1 (C.elegans)-like 3 (myoferlin) (FER1L3), mRNA. /FEA=mRNA /GEN=FER1L3 /PROD=fer-1 (C.elegans)-like 3 (myoferlin) /DB\_XREF=gi:7305052 /UG=Hs.234680 fer-1 (C.elegans)-like 3 (myoferlin) /FL=gb:AF182316.1 gb:NM\_013451.1

Affy Probe_ID	Sequence Description
205680_at	gb:NM_002425.1 /DEF=Homo sapiens matrix metalloproteinase 10 (stromelysin 2) (MMP10), mRNA. /FEA=mRNA /GEN=MMP10 /PROD=matrix metalloproteinase 10 preproprotein /DB_XREF=gi:4505204 /UG=Hs.2258 matrix metalloproteinase 10 (stromelysin 2) /FL=gb:BC002591.1 gb:NM_002425.1
214316_x_at	Consensus includes gb:AI378706 /FEA=EST /DB_XREF=gi:4188559 /DB_XREF=est:tb91f09.x1 /CLONE=IMAGE:2061737 /UG=Hs.16488 calreticulin
212919_at	Consensus includes gb:AV715578 /FEA=EST /DB_XREF=gi:10797095 /DB_XREF=est:AV715578 /CLONE=DCBBJG09 /UG=Hs.21056 Homo sapiens cDNA: FLJ21366 fis, clone COL03012, highly similar to AB002445 Homo sapiens mRNA from chromosome 5q21-22
203917_at	gb:NM_001338.1 /DEF=Homo sapiens coxsackie virus and adenovirus receptor (CXADR), mRNA. /FEA=mRNA /GEN=CXADR /PROD=coxsackie virus and adenovirus receptor /DB_XREF=gi:4503172 /UG=Hs.79187 coxsackie virus and adenovirus receptor /FL=gb:BC003684.1 gb:U90716.1 gb:NM_001338.1
214014_at	Consensus includes gb:W81196 /FEA=EST /DB_XREF=gi:1392235 /DB_XREF=est:zd85d03.s1 /CLONE=IMAGE:347429 /UG=Hs.12289 Cdc42 effector protein 2
219051_x_at	gb:NM_024042.1 /DEF=Homo sapiens hypothetical protein MGC2601 (MGC2601), mRNA. /FEA=mRNA /GEN=MGC2601 /PROD=hypothetical protein MGC2601 /DB_XREF=gi:13128999 /UG=Hs.124915 hypothetical protein MGC2601 /FL=gb:BC000662.1 gb:NM_024042.1
219135_s_at	Consensus includes gb:AL515916 /FEA=EST /DB_XREF=gi:12779409 /DB_XREF=est:AL515916 /CLONE=CS0DA001YH18 (3 prime) /UG=Hs.58362 hypothetical protein FLJ12681 /FL=gb:NM_022773.1
211143_x_at	gb:D49728.1 /DEF=Human NAK1 mRNA for DNA binding protein, complete cds. /FEA=mRNA /GEN=NAK1 /PROD=DNA binding protein /DB_XREF=gi:1813881 /UG=Hs.1119 nuclear receptor subfamily 4, group A, member 1 /FL=gb:D49728.1
203188_at	gb:NM_006876.1 /DEF=Homo sapiens i-beta-1,3-N-acetylglucosaminyltransferase (BETA3GNTI), mRNA. /FEA=mRNA /GEN=BETA3GNTI /PROD=i-beta-1,3-N-acetylglucosaminyltransferase /DB_XREF=gi:5802983 /UG=Hs.8526 i-beta-1,3-N-acetylglucosaminyltransferase /FL=gb:AF029893.1 gb:NM_006876.1
205839_s_at	gb:NM_004758.1 /DEF=Homo sapiens peripheral benzodiazepine receptor-associated protein 1 (PRAX-1), mRNA. /FEA=mRNA /GEN=PRAX-1 /PROD=peripheral benzodiazepine receptor-associated protein 1 /DB_XREF=gi:4758955 /UG=Hs.112499 peripheral benzodiazepine receptor-associated protein 1 /FL=gb:AF039571.1 gb:NM_004758.1

**Affy Probe ID**

**Sequence Description**

200890_s_at	Consensus includes gb:AW006345 /FEA=EST /DB_XREF=gi:5855123 /DB_XREF=est:wt04d05.x1 /CLONE=IMAGE:2506473 /UG=Hs.250773 signal sequence receptor, alpha (translocon-associated protein alpha) /FL=gb:AF156965.1 gb:NM_003144.2
202967_at	gb:NM_001512.1 /DEF=Homo sapiens glutathione S-transferase A4 (GSTA4), mRNA. /FEA=mRNA /GEN=GSTA4 /PROD=glutathione S-transferase A4 /DB_XREF=gi:4504172 /UG=Hs.169907 glutathione S-transferase A4 /FL=gb:AF020918.1 gb:AF025887.1 gb:NM_001512.1 gb:AF125271.1 gb:AF125272.1 gb:AF125273.1
37170_at	Cluster Incl. AB015331:Homo sapiens HRIHFB2017 mRNA, partial cds /cds=(0,650) /gb=AB015331 /gi=3970851 /ug=Hs.108903 /len=1064
201830_s_at	gb:NM_005863.1 /DEF=Homo sapiens neuroepithelial cell transforming gene 1 (NET1), mRNA. /FEA=mRNA /GEN=NET1 /PROD=neuroepithelial cell transforming gene 1 /DB_XREF=gi:5031938 /UG=Hs.25155 neuroepithelial cell transforming gene 1 /FL=gb:NM_005863.1 gb:U02081.1
213825_at	Consensus includes gb:AA757419 /FEA=EST /DB_XREF=gi:2805282 /DB_XREF=est:zg36b11.s1 /CLONE=IMAGE:395421 /UG=Hs.176977 protein kinase C binding protein 2
207540_s_at	gb:NM_003177.1 /DEF=Homo sapiens spleen tyrosine kinase (SYK), mRNA. /FEA=mRNA /GEN=SYK /PROD=spleen tyrosine kinase /DB_XREF=gi:4507328 /UG=Hs.74101 spleen tyrosine kinase /FL=gb:NM_003177.1
207008_at	gb:NM_001557.1 /DEF=Homo sapiens interleukin 8 receptor, beta (IL8RB), mRNA. /FEA=mRNA /GEN=IL8RB /PROD=interleukin 8 receptor, beta /DB_XREF=gi:4504682 /UG=Hs.846 interleukin 8 receptor, beta /FL=gb:M94582.1 gb:M73969.1 gb:NM_001557.1 gb:L19593.1
212671_s_at	Consensus includes gb:BG397856 /FEA=EST /DB_XREF=gi:13291304 /DB_XREF=est:602438950F1 /CLONE=IMAGE:4564956 /UG=Hs.198253 major histocompatibility complex, class II, DQ alpha 1
200962_at	Consensus includes gb:AI348010 /FEA=EST /DB_XREF=gi:4085216 /DB_XREF=est:qp61b10.x1 /CLONE=IMAGE:1927483 /UG=Hs.184014 ribosomal protein L31 /FL=gb:NM_000993.1
205379_at	gb:NM_001236.2 /DEF=Homo sapiens carbonyl reductase 3 (CBR3), mRNA. /FEA=mRNA /GEN=CBR3 /PROD=carbonyl reductase 3 /DB_XREF=gi:7108334 /UG=Hs.154510 carbonyl reductase 3 /FL=gb:BC002812.1 gb:AB004854.1 gb:NM_001236.2
205599_at	gb:NM_005658.1 /DEF=Homo sapiens TNF receptor-associated factor 1 (TRAF1), mRNA. /FEA=mRNA /GEN=TRAF1 /PROD=TNF receptor-associated factor 1 /DB_XREF=gi:5032192 /UG=Hs.2134 TNF receptor-associated factor 1 /FL=gb:NM_005658.1 gb:U19261.1

Affy Probe_ID	Sequence Description
206382_s_at	gb:NM_001709.1 /DEF=Homo sapiens brain-derived neurotrophic factor (BDNF), mRNA. /FEA=mRNA /GEN=BDNF /PROD=brain-derived neurotrophic factor /DB_XREF=gi:4502392 /UG=Hs.56023 brain-derived neurotrophic factor /FL=gb:NM_001709.1
215097_at	Consensus includes gb:BG420053 /FEA=EST /DB_XREF=gi:13326559 /DB_XREF=est:602453740F1 /CLONE=IMAGE:4592120 /UG=Hs.76368 capping protein (actin filament) muscle Z-line, beta
205130_at	gb:NM_014226.1 /DEF=Homo sapiens renal tumor antigen (RAGE), mRNA. /FEA=mRNA /GEN=RAGE /PROD=renal tumor antigen /DB_XREF=gi:7657497 /UG=Hs.104119 renal tumor antigen /FL=gb:AB022694.1 gb:NM_014226.1
217497_at	Consensus includes gb:AW613387 /FEA=EST /DB_XREF=gi:7318573 /DB_XREF=est:hh71e04.x1 /CLONE=IMAGE:2968254 /UG=Hs.278541 ESTs, Moderately similar to TYPH_HUMAN THYMIDINE PHOSPHORYLASE PRECURSOR H.sapiens
221589_s_at	Consensus includes gb:AW612403 /FEA=EST /DB_XREF=gi:7317589 /DB_XREF=est:hh01b05.x1 /CLONE=IMAGE:2953809 /UG=Hs.293970 methylmalonate-semialdehyde dehydrogenase /FL=gb:AF130089.1
213437_at	Consensus includes gb:AA861784 /FEA=EST /DB_XREF=gi:2953924 /DB_XREF=est:ak35d05.s1 /CLONE=IMAGE:1407945 /UG=Hs.7972 KIAA0871 protein
210891_s_at	gb:AF035737.1 /DEF=Homo sapiens general transcription factor 2-1 (GTF2I) mRNA, complete cds. /FEA=mRNA /GEN=GTF2I /PROD=general transcription factor 2-I; alternativelyspliced product /DB_XREF=gi:2827179 /UG=Hs.278589 general transcription factor II, i /FL=gb:AF035737.1
216942_s_at	Consensus includes gb:D28586.1 /DEF=Human mRNA for LFA-3(delta D2), partial cds. /FEA=mRNA /PROD=LFA-3(delta D2) /DB_XREF=gi:466540 /UG=Hs.75626 CD58 antigen, (lymphocyte function-associated antigen 3)
203154_s_at	gb:NM_005884.2 /DEF=Homo sapiens protein kinase related to S. cerevisiae STE20, effector for Cdc42Hs (PAK4), mRNA. /FEA=mRNA /GEN=PAK4 /PROD=protein kinase related to S. cerevisiae STE20, effector for Cdc42Hs /DB_XREF=gi:7382497 /UG=Hs.20447 protein kinase related to S. cerevisiae STE20, effector for Cdc42Hs /FL=gb:AF005046.1 gb:NM_005884.2
218852_at	gb:NM_017917.1 /DEF=Homo sapiens hypothetical protein FLJ20644 (FLJ20644), mRNA. /FEA=mRNA /GEN=FLJ20644 /PROD=hypothetical protein FLJ20644 /DB_XREF=gi:8923599 /UG=Hs.49376 hypothetical protein FLJ20644 /FL=gb:NM_017917.1

Affy Probe_ID	Sequence Description
201315_x_at	gb:NM_006435.1 /DEF=Homo sapiens interferon induced transmembrane protein 2 (1-8D) (IFITM2), mRNA. /FEA=mRNA /GEN=IFITM2 /PROD=interferon induced transmembrane protein 2(1-8D) /DB_XREF=gi:10835237 /UG=Hs.174195 interferon induced transmembrane protein 2 (1-8D) /FL=gb:NM_006435.1
46665_at	Cluster Incl. AI949392:wq11a04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2470926 /clone_end=3 /gb=AI949392 /gi=5741790 /ug=Hs.7188 /len=596
201616_s_at	Consensus includes gb:AL577531 /FEA=EST /DB_XREF=gi:12940753 /DB_XREF=est:AL577531 /CLONE=CS0DI087YP20 (3 prime) /UG=Hs.325474 caldesmon 1 /FL=gb:NM_004342.2 gb:M64110.1
219165_at	gb:NM_021630.1 /DEF=Homo sapiens PDZ-LIM protein mystique (LOC59346), mRNA. /FEA=mRNA /GEN=LOC59346 /PROD=PDZ-LIM protein mystique /DB_XREF=gi:11055999 /UG=Hs.19447 PDZ-LIM protein mystique /FL=gb:AY007729.1 gb:NM_021630.1
59625_at	Cluster Incl. AI912351:tz24d02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2289507 /clone_end=3 /gb=AI912351 /gi=5632206 /ug=Hs.198793 /len=507
216037_x_at	Consensus includes gb:AA664011 /FEA=EST /DB_XREF=gi:2618002 /DB_XREF=est:ac03f10.s1 /CLONE=IMAGE:855403 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG-box)
210874_s_at	gb:BC004483.1 /DEF=Homo sapiens, putative tumor suppressor, clone MGC:10586, mRNA, complete cds. /FEA=mRNA /PROD=putative tumor suppressor /DB_XREF=gi:13325345 /UG=Hs.22919 putative tumor suppressor /FL=gb:BC004483.1
201829_at	Consensus includes gb:AW263232 /FEA=EST /DB_XREF=gi:6640048 /DB_XREF=est:xn79e03.x1 /CLONE=IMAGE:2700700 /UG=Hs.25155 neuroepithelial cell transforming gene 1 /FL=gb:NM_005863.1 gb:U02081.1
202627_s_at	Consensus includes gb:AL574210 /FEA=EST /DB_XREF=gi:12934198 /DB_XREF=est:AL574210 /CLONE=CS0DI039YB02 (3 prime) /UG=Hs.82085 serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 /FL=gb:NM_000602.1 gb:M16006.1
202902_s_at	gb:NM_004079.1 /DEF=Homo sapiens cathepsin S (CTSS), mRNA. /FEA=mRNA /GEN=CTSS /PROD=cathepsin S /DB_XREF=gi:4758097 /UG=Hs.181301 cathepsin S /FL=gb:BC002642.1 gb:M86553.1 gb:NM_004079.1 gb:M90696.1
218456_at	gb:NM_023925.1 /DEF=Homo sapiens hypothetical protein FLJ22569 (FLJ22569), mRNA. /FEA=mRNA /GEN=FLJ22569 /PROD=hypothetical protein FLJ22569 /DB_XREF=gi:12965192 /UG=Hs.234355 hypothetical protein FLJ22569 /FL=gb:NM_023925.1

Affy Probe_ID	Sequence Description
217529_at	Consensus includes gb:BE547674 /FEA=EST /DB_XREF=gi:9776319 /DB_XREF=est:601076255F1 /CLONE=IMAGE:3462325 /UG=Hs.204169 ESTs
200833_s_at	gb:NM_015646.1 /DEF=Homo sapiens RAP1B, member of RAS oncogene family (RAP1B), mRNA. /FEA=mRNA /GEN=RAP1B /PROD=DKFZP586H0723 protein /DB_XREF=gi:7661677 /UG=Hs.156764 RAP1B, member of RAS oncogene family /FL=gb:BC000176.2 gb:NM_015646.1
218217_at	gb:NM_021626.1 /DEF=Homo sapiens serine carboxypeptidase 1 precursor protein (HSCP1), mRNA. /FEA=mRNA /GEN=HSCP1 /PROD=serine carboxypeptidase 1 precursor protein /DB_XREF=gi:11055991 /UG=Hs.106747 serine carboxypeptidase 1 precursor protein /FL=gb:AF282618.1 gb:NM_021626.1 gb:AF113214.1 gb:AF265441.1
209493_at	gb:AF338650.1 /DEF=Homo sapiens PDZ domain-containing protein AIPC (AIPC) mRNA, complete cds. /FEA=mRNA /GEN=AIPC /PROD=PDZ domain-containing protein AIPC /DB_XREF=gi:12751451 /UG=Hs.173035 KIAA0300 protein /FL=gb:AF338650.1
202937_x_at	Consensus includes gb:AL022316 /DEF=Human DNA sequence from clone CTA-126B4 on chromosome 22q13.2-13.31 Contains two or three novel genes, ESTs, STSs, GSSs and a CpG Island /FEA=mRNA_3 /DB_XREF=gi:4691242 /UG=Hs.239934 CGI-96 protein /FL=gb:AF151854.1 gb:NM_015703.1
207055_at	gb:NM_004767.1 /DEF=Homo sapiens endothelin type b receptor-like protein 2 (ET(B)R-LP-2), mRNA. /FEA=mRNA /GEN=ET(B)R-LP-2 /PROD=endothelin type b receptor-like protein 2 /DB_XREF=gi:4758309 /UG=Hs.132049 endothelin type b receptor-like protein 2 /FL=gb:NM_004767.1
212731_at	Consensus includes gb:U79297.1 /DEF=Human clone 23589 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710280 /UG=Hs.11506 Human clone 23589 mRNA sequence
201772_at	gb:NM_015878.1 /DEF=Homo sapiens antizyme inhibitor (LOC51582), mRNA. /FEA=mRNA /GEN=LOC51582 /PROD=antizyme inhibitor /DB_XREF=gi:7706219 /UG=Hs.223014 antizyme inhibitor /FL=gb:D88674.1 gb:NM_015878.1
210871_x_at	gb:AL133046.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C0515 (from clone DKFZp434C0515); complete cds. /FEA=mRNA /GEN=DKFZp434C0515 /PROD=hypothetical protein /DB_XREF=gi:6453458 /UG=Hs.22587 KIAA0923 protein /FL=gb:AL133046.1
212386_at	Consensus includes gb:BF592782 /FEA=EST /DB_XREF=gi:11685106 /DB_XREF=est:7j94d06.x1 /CLONE=IMAGE:3442594 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1000272

**Affy Probe ID**

**Sequence Description**

202575\_at

gb:NM\_001878.2 /DEF=Homo sapiens cellular retinoic acid-binding protein 2 (CRABP2), mRNA. /FEA=mRNA /GEN=CRABP2 /PROD=cellular retinoic acid-binding protein 2 /DB\_XREF=gi:6382069 /UG=Hs.183650 cellular retinoic acid-binding protein 2 /FL=gb:BC001109.1 gb:M68867.1 gb:NM\_001878.2

219388\_at

gb:NM\_024915.1 /DEF=Homo sapiens hypothetical protein FLJ13782 (FLJ13782), mRNA. /FEA=mRNA /GEN=FLJ13782 /PROD=hypothetical protein FLJ13782 /DB\_XREF=gi:13376381 /UG=Hs.257924 hypothetical protein FLJ13782 /FL=gb:NM\_024915.1

218910\_at

gb:NM\_018075.1 /DEF=Homo sapiens hypothetical protein FLJ10375 (FLJ10375), mRNA. /FEA=mRNA /GEN=FLJ10375 /PROD=hypothetical protein FLJ10375 /DB\_XREF=gi:8922383 /UG=Hs.319088 hypothetical protein FLJ10375 /FL=gb:NM\_018075.1

202308\_at

gb:NM\_004176.1 /DEF=Homo sapiens sterol regulatory element binding transcription factor 1 (SREBF1), mRNA. /FEA=mRNA /GEN=SREBF1 /PROD=sterol regulatory element binding transcriptionfactor 1 /DB\_XREF=gi:4759167 /UG=Hs.166 sterol regulatory element binding transcription factor 1 /FL=gb:U00968.1 gb:NM\_004176.1

222146\_s\_at

Consensus includes gb:AK026674.1 /DEF=Homo sapiens cDNA: FLJ23021 fis, clone LNG01014, highly similar to HUMSEF21B Human SEF2-1B protein (SEF2-1B) mRNA. /FEA=mRNA /DB\_XREF=gi:10439577 /UG=Hs.326198 transcription factor 4

212953\_x\_at

Consensus includes gb:BE251303 /FEA=EST /DB\_XREF=gi:9121413 /DB\_XREF=est:601107814F1 /CLONE=IMAGE:3344018 /UG=Hs.16488 calreticulin

213306\_at

Consensus includes gb:AA917899 /FEA=EST /DB\_XREF=gi:3057789 /DB\_XREF=est:ol76e05.s1 /CLONE=IMAGE:1535552 /UG=Hs.169378 multiple PDZ domain protein

213309\_at

Consensus includes gb:AL117515.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L0735 (from clone DKFZp434L0735); partial cds. /FEA=mRNA /GEN=DKFZp434L0735 /PROD=hypothetical protein /DB\_XREF=gi:5912029 /UG=Hs.54886 phospholipase C, epsilon 2

204115\_at

gb:NM\_004126.1 /DEF=Homo sapiens guanine nucleotide binding protein 11 (GNG11), mRNA. /FEA=mRNA /GEN=GNG11 /PROD=guanine nucleotide binding protein 11 /DB\_XREF=gi:4758447 /UG=Hs.83381 guanine nucleotide binding protein 11 /FL=gb:NM\_004126.1 gb:U31384.1

212850\_s\_at

Consensus includes gb:AA584297 /FEA=EST /DB\_XREF=gi:2368906 /DB\_XREF=est:nn79e08.s1 /CLONE=IMAGE:1090118 /UG=Hs.4930 low density lipoprotein receptor-related protein 4

218943\_s\_at

gb:NM\_014314.1 /DEF=Homo sapiens RNA helicase (RIG-I), mRNA. /FEA=mRNA /GEN=RIG-I /PROD=RNA helicase /DB\_XREF=gi:7657515 /UG=Hs.145612 RNA helicase /FL=gb:AF038963.1 gb:NM\_014314.1



# Affy Probe\_ID

## Sequence Description

210119\_at gb:U73191.1 /DEF=Human inward rectifier potassium channel (Kir1.3), complete cds. /FEA=mRNA /GEN=Kir1.3 /PROD=inward rectifier potassium channel /DB\_XREF=gi:1765984 /UG=Hs.17287 potassium inwardly-rectifying channel, subfamily J, member 15 /FL=gb:U73191.1 gb:NM\_002243.1

202731\_at gb:NM\_014456.1 /DEF=Homo sapiens programmed cell death 4 (PDCD4), mRNA. /FEA=mRNA /GEN=PDCD4 /PROD=programmed cell death 4 /DB\_XREF=gi:7657448 /UG=Hs.296251 programmed cell death 4 /FL=gb:U96628.1 gb:NM\_014456.1

212573\_at Consensus includes gb:AF131747.1 /DEF=Homo sapiens clone 24951 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:4406562 /UG=Hs.167115 KIAA0830 protein

218999\_at gb:NM\_018295.1 /DEF=Homo sapiens hypothetical protein FLJ11000 (FLJ11000), mRNA. /FEA=mRNA /GEN=FLJ11000 /PROD=hypothetical protein FLJ11000 /DB\_XREF=gi:8922813 /UG=Hs.302981 hypothetical protein FLJ11000 /FL=gb:NM\_018295.1

212095\_s\_at Consensus includes gb:BE552421 /FEA=EST /DB\_XREF=gi:9794113 /DB\_XREF=est:hw26b02.x1 /CLONE=IMAGE:3184011 /UG=Hs.7946 KIAA1288 protein

203608\_at Consensus includes gb:AL031230 /DEF=Human DNA sequence from clone 73M23 on chromosome 6p22.2-22.3. Contains the 5 part of the possibly alternatively spliced gene for Phosphatidylinositol-glycan-specific Phospholipase D 1 precursor (EC 3.1.4.50, PIGPLD1, Glycoprotein Phospholipase D... /FEA=mRNA\_5 /DB\_XREF=gi:3947845 /UG=Hs.5299 aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase) /FL=gb:NM\_001080.1

202733\_at gb:NM\_004199.1 /DEF=Homo sapiens procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II (P4HA2), mRNA. /FEA=mRNA /GEN=P4HA2 /PROD=procollagen-proline, 2-oxoglutarate4-dioxygenase (proline 4-hydroxylase), alpha polypeptideII /DB\_XREF=gi:4758867 /UG=Hs.3622 procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II /FL=gb:U90441.1 gb:NM\_004199.1

218087\_s\_at gb:NM\_015385.1 /DEF=Homo sapiens SH3-domain protein 5 (ponsin) (SH3D5), mRNA. /FEA=mRNA /GEN=SH3D5 /PROD=ponsin /DB\_XREF=gi:7661699 /UG=Hs.108924 SH3-domain protein 5 (ponsin) /FL=gb:AL117472.1 gb:NM\_015385.1

217047\_s\_at Consensus includes gb:AK027138.1 /DEF=Homo sapiens cDNA: FLJ23485 fis, clone KAIJA05211. /FEA=mRNA /DB\_XREF=gi:10440190 /UG=Hs.177664 KIAA0914 gene product

# Affy Probe ID

## Sequence Description

220524\_at gb:NM\_024823.1 /DEF=Homo sapiens hypothetical protein FLJ21596 (FLJ21596), mRNA. /FEA=mRNA /GEN=FLJ21596 /PROD=hypothetical protein FLJ21596 /DB\_XREF=gi:13376219 /UG=Hs.171537 hypothetical protein FLJ21596 /FL=gb:NM\_024823.1

202872\_at Consensus includes gb:AW024925 /FEA=EST /DB\_XREF=gi:5878455 /DB\_XREF=est:wu92g09.x1 /CLONE=IMAGE:2527552 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM\_001695.1

209979\_at gb:AF001042.1 /DEF=Homo sapiens RNA editase (RED1) mRNA, complete cds. /FEA=mRNA /GEN=RED1 /PROD=RNA editase /DB\_XREF=gi:2114492 /UG=Hs.85302 adenosine deaminase, RNA-specific, B1 (homolog of rat RED1) /FL=gb:AF001042.1

205651\_x\_at gb:NM\_007023.1 /DEF=Homo sapiens cAMP-regulated guanine nucleotide exchange factor II (CAMP-GEFII), mRNA. /FEA=mRNA /GEN=CAMP-GEFII /PROD=cAMP-regulated guanine nucleotide exchange factor II /DB\_XREF=gi:5901913 /UG=Hs.91971 cAMP-regulated guanine nucleotide exchange factor II /FL=gb:U78516.1

202531\_at gb:NM\_007023.1

200731\_s\_at gb:NM\_002198.1 /DEF=Homo sapiens interferon regulatory factor 1 (IRF1), mRNA. /FEA=mRNA /GEN=IRF1 /PROD=interferon regulatory factor 1 /DB\_XREF=gi:4504720 /UG=Hs.80645 interferon regulatory factor 1 /FL=gb:NM\_002198.1

200731\_s\_at Consensus includes gb:AW165960 /FEA=EST /DB\_XREF=gi:6397485 /DB\_XREF=est:xf43a12.x1 /CLONE=IMAGE:2620798 /UG=Hs.227777 protein tyrosine phosphatase type IVA, member 1 /FL=gb:U48296.1

213058\_at gb:NM\_003463.1

213058\_at Consensus includes gb:AL033538 /DEF=Human DNA sequence from clone RP3-477H23 on chromosome 22q12.1-12.2 Contains parts of one or two novel genes, ESTs, STSs and GSSs /FEA=mRNA /DB\_XREF=gi:4165236 /UG=Hs.11390 KIAA1043 protein

215253\_s\_at Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586D0518 (from clone DKFZp586D0518). /FEA=mRNA /DB\_XREF=gi:4500160 /UG=Hs.250724 Homo sapiens mRNA; cDNA DKFZp586D0518 (from clone DKFZp586D0518)

200628\_s\_at gb:M61715.1 /DEF=Human tryptophanyl-tRNA synthetase (WRS) mRNA, complete cds. /FEA=mRNA /GEN=WARS /PROD=transfer RNA-Trp synthetase /DB\_XREF=gi:340367 /UG=Hs.82030 tryptophanyl-tRNA synthetase /FL=gb:M77804.1 gb:M61715.1 gb:NM\_004184.2

# Affy Probe ID

## Sequence Description

218638_s_at	gb:NM_012445.1 /DEF=Homo sapiens spondin 2, extracellular matrix protein (SPON2), mRNA. /FEA=mRNA /GEN=SPON2 /PROD=spondin 2, extracellular matrix protein /DB_XREF=gi:6912681 /UG=Hs.288126 spondin 2, extracellular matrix protein /FL=gb:BC002707.1 gb:AB027466.1 gb:NM_012445.1
206393_at	gb:NM_003282.1 /DEF=Homo sapiens troponin I, skeletal, fast (TNNI2), mRNA. /FEA=mRNA /GEN=TNNI2 /PROD=troponin I, skeletal, fast /DB_XREF=gi:4507620 /UG=Hs.83760 troponin I, skeletal, fast /FL=gb:NM_003282.1 gb:L21715.1
218062_x_at	gb:NM_012121.2 /DEF=Homo sapiens Cdc42 effector protein 4; binder of Rho GTPases 4 (CEP4), mRNA. /FEA=mRNA /GEN=CEP4 /PROD=Cdc42 effector protein 4; binder of Rho GTPases4 /DB_XREF=gi:13786126 /UG=Hs.3903 Cdc42 effector protein 4; binder of Rho GTPases 4 /FL=gb:AB042237.1 gb:NM_012121.2 gb:AF099664.1
203298_s_at	gb:NM_004973.2 /DEF=Homo sapiens jumonji (mouse) homolog (JMJ), mRNA. /FEA=mRNA /GEN=JMJ /PROD=jumonji /DB_XREF=gi:11863151 /UG=Hs.40154 jumonji (mouse) homolog /FL=gb:NM_004973.2 gb:U57592.1
213324_at	Consensus includes gb:AK024281.1 /DEF=Homo sapiens cDNA FLJ14219 fis, clone NT2RP3003800, highly similar to Rattus norvegicus tyrosine protein kinase pp60-c-src mRNA. /FEA=mRNA /DB_XREF=gi:10436622 /UG=Hs.198298 v-src avian sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog
213591_at	Consensus includes gb:AU149534 /FEA=EST /DB_XREF=gi:11011055 /DB_XREF=est:AU149534 /CLONE=NT2RM4002512 /UG=Hs.74294 aldehyde dehydrogenase 7 family, member A1
205047_s_at	gb:NM_001673.1 /DEF=Homo sapiens asparagine synthetase (ASNS), mRNA. /FEA=mRNA /GEN=ASNS /PROD=asparagine synthetase /DB_XREF=gi:4502258 /UG=Hs.75692 asparagine synthetase /FL=gb:M27396.1 gb:NM_001673.1
220066_at	gb:NM_022162.1 /DEF=Homo sapiens NOD2 protein (NOD2), mRNA. /FEA=mRNA /GEN=NOD2 /PROD=NOD2 protein /DB_XREF=gi:11545911 /UG=Hs.135201 NOD2 protein /FL=gb:AF178930.1 gb:NM_022162.1
217906_at	gb:NM_014315.1 /DEF=Homo sapiens host cell factor homolog (LCP), mRNA. /FEA=mRNA /GEN=LCP /PROD=host cell factor homolog /DB_XREF=gi:7657300 /UG=Hs.20597 host cell factor homolog /FL=gb:BC002335.1 gb:AF113131.1 gb:NM_014315.1 gb:AF244137.1
202023_at	gb:NM_004428.1 /DEF=Homo sapiens ephrin-A1 (EFNA1), mRNA. /FEA=mRNA /GEN=EFNA1 /PROD=ephrin A1 precursor /DB_XREF=gi:4758245 /UG=Hs.1624 ephrin-A1 /FL=gb:M57730.1 gb:NM_004428.1

Affy Probe_ID	Sequence Description
221751_at	Consensus includes gb:AL565516 /FEA=EST /DB_XREF=gi:12916970 /DB_XREF=est:AL565516 /CLONE=CS0DF004YF08 (3 prime) /UG=Hs.6986 Human glucose transporter pseudogene
201463_s_at	gb:Nm_006755.1 /DEF=Homo sapiens transaldolase 1 (TALDO1), mRNA. /FEA=mRNA /GEN=TALDO1 /PROD=transaldolase 1 /DB_XREF=gi:5803186 /UG=Hs.77290 transaldolase 1 /FL=gb:U19437.2 gb:Nm_006755.1
203763_at	gb:Nm_016008.1 /DEF=Homo sapiens CGI-60 protein (LOC51626), mRNA. /FEA=mRNA /GEN=LOC51626 /PROD=CGI-60 protein /DB_XREF=gi:7706299 /UG=Hs.7627 CGI-60 protein /FL=gb:AF151818.1 gb:Nm_016008.1
205383_s_at	gb:Nm_015642.1 /DEF=Homo sapiens zinc finger protein 288 (ZNF288), mRNA. /FEA=mRNA /GEN=ZNF288 /PROD=zinc finger protein 288 /DB_XREF=gi:7661651 /UG=Hs.159456 zinc finger protein 288 /FL=gb:AL050276.1 gb:Nm_015642.1
213373_s_at	Consensus includes gb:BF439983 /FEA=EST /DB_XREF=gi:11452500 /DB_XREF=est:nac51f11.x1 /CLONE=IMAGE:3406220 /UG=Hs.19949 caspase 8, apoptosis-related cysteine protease /FL=gb:U58143.1
200707_at	gb:Nm_002743.1 /DEF=Homo sapiens protein kinase C substrate 80K-H (PRKCSH), mRNA. /FEA=mRNA /GEN=PRKCSH /PROD=protein kinase C substrate 80K-H /DB_XREF=gi:4506076 /UG=Hs.1432 protein kinase C substrate 80K-H /FL=gb:J03075.1 gb:Nm_002743.1 gb:AF144075.1
212427_at	Consensus includes gb:AB002366.1 /DEF=Human mRNA for KIAA0368 gene, partial cds. /FEA=mRNA /GEN=KIAA0368 /DB_XREF=gi:2224676 /UG=Hs.3852 KIAA0368 protein
216268_s_at	Consensus includes gb:U77914.1 /DEF=Human soluble protein Jagged mRNA, partial cds. /FEA=mRNA /PROD=soluble protein Jagged /DB_XREF=gi:1684889 /UG=Hs.91143 jagged 1 (Alagille syndrome)
210457_x_at	gb:AF176039.1 /DEF=Homo sapiens high mobility group protein-R mRNA, complete cds. /FEA=mRNA /PROD=high mobility group protein-R /DB_XREF=gi:5834272 /UG=Hs.139800 high-mobility group (nonhistone chromosomal) protein isoforms I and Y /FL=gb:AF176039.1
200924_s_at	gb:Nm_002394.1 /DEF=Homo sapiens solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 (SLC3A2), mRNA. /FEA=mRNA /GEN=SLC3A2 /PROD=antigen identified by monoclonal antibodies 4F2, TRA1.10, TROP4, and T43 /DB_XREF=gi:4505140 /UG=Hs.79748 solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2 /FL=gb:BC001061.1 gb:J02769.1 gb:J03569.1 gb:Nm_002394.1 gb:AB018010.1

Affy Probe_ID	Sequence Description
203409_at	gb:NM_000107.1 /DEF=Homo sapiens damage-specific DNA binding protein 2 (48kD) (DDB2), mRNA. /FEA=mRNA /GEN=DDB2 /PROD=damage-specific DNA binding protein 2 (48kD) /DB_XREF=gi:4557514 /UG=Hs.77602 damage-specific DNA binding protein 2 (48kD) /FL=gb:BC000093.1 gb:U18300.1 gb:NM_000107.1 gb:NM_004334.1 /DEF=Homo sapiens bone marrow stromal cell antigen 1 (BST1), mRNA. /FEA=mRNA /GEN=BST1 /PROD=bone marrow stromal cell antigen 1 precursor /DB_XREF=gi:4757873 /UG=Hs.169998 bone marrow stromal cell antigen 1 /FL=gb:NM_004334.1 gb:D21878.1
213609_s_at	Consensus includes gb:AB023144.1 /DEF=Homo sapiens mRNA for KIAA0927 protein, partial cds. /FEA=mRNA /GEN=KIAA0927 /PROD=KIAA0927 protein /DB_XREF=gi:4589485 /UG=Hs.194766 seizure related gene 6 (mouse)-like
212932_at	Consensus includes gb:AK022494.1 /DEF=Homo sapiens cDNA FLJ12432 fis, clone NT2RM1000018, highly similar to Human mRNA for KIAA0066 gene. /FEA=mRNA /DB_XREF=gi:10433912 /UG=Hs.227881 RAB3 GTPase-ACTIVATING PROTEIN
204632_at	gb:NM_003942.1 /DEF=Homo sapiens ribosomal protein S6 kinase, 90kD, polypeptide 4 (RPS6KA4), mRNA. /FEA=mRNA /GEN=RPS6KA4 /PROD=ribosomal protein S6 kinase, 90kD, polypeptide4 /DB_XREF=gi:4506734 /UG=Hs.105584 ribosomal protein S6 kinase, 90kD, polypeptide 4 /FL=gb:NM_003942.1
201988_s_at	Consensus includes gb:BF438056 /FEA=EST /DB_XREF=gi:11450573 /DB_XREF=est:7q66e01.x1 /CLONE=IMAGE:3703369 /UG=Hs.13313 cAMP responsive element binding protein-like 2 /FL=gb:AF039081.1 gb:NM_001310.1
212117_at	Consensus includes gb:BF978689 /FEA=EST /DB_XREF=gi:12345904 /DB_XREF=est:602149457F2 /CLONE=IMAGE:4308014 /UG=Hs.166982 phosphatidylinositol glycan, class F
217496_s_at	Consensus includes gb:AA918442 /FEA=EST /DB_XREF=gi:3058332 /DB_XREF=est:om38b01.s1 /CLONE=IMAGE:1543273 /UG=Hs.326641 Homo sapiens insulin-degrading enzyme (IDE), mRNA
203180_at	gb:NM_000693.1 /DEF=Homo sapiens aldehyde dehydrogenase 1 family, member A3 (ALDH1A3), mRNA. /FEA=mRNA /GEN=ALDH1A3 /PROD=aldehyde dehydrogenase 1A3 /DB_XREF=gi:4502040 /UG=Hs.75746 aldehyde dehydrogenase 1 family, member A3 /FL=gb:NM_000693.1 gb:U07919.1
203801_at	Consensus includes gb:AA013164 /FEA=EST /DB_XREF=gi:1474190 /DB_XREF=est:ze35e12.s1 /CLONE=IMAGE:361006 /UG=Hs.247324 hypothetical protein LOC63931 /FL=gb:NM_022100.1

# Affy Probe\_ID

## Sequence Description

214494\_s\_at Consensus includes gb:NM\_005200.1 /DEF=Homo sapiens cell matrix adhesion regulator (CMAR), mRNA. /FEA=CDS /GEN=CMAR /PROD=cell matrix adhesion regulator /DB\_XREF=gi:4885142 /UG=Hs.272480 cell matrix adhesion regulator /FL=gb:NM\_005200.1

201196\_s\_at gb:M21154.1 /DEF=Human S-adenosylmethionine decarboxylase mRNA, complete cds. /FEA=mRNA /GEN=AMD2 /DB\_XREF=gi:178517 /UG=Hs.262476 S-adenosylmethionine decarboxylase 1 /FL=gb:BC000171.2 gb:M21154.1 gb:NM\_001634.3

210325\_at gb:M28825.1 /DEF=Human thymocyte antigen CD1a mRNA, complete cds. /FEA=mRNA /GEN=CD1A /DB\_XREF=gi:180035 /UG=Hs.1309 CD1A antigen, a polypeptide /FL=gb:M28825.1

200755\_s\_at Consensus includes gb:BF939365 /FEA=EST /DB\_XREF=gi:12356685 /DB\_XREF=est:nad87h04.x1 /CLONE=IMAGE:3410551 /UG=Hs.7753 calumenin /FL=gb:U67280.1 gb:AF013759.1 gb:NM\_001219.2

206580\_s\_at gb:NM\_016938.1 /DEF=Homo sapiens EGF-containing fibulin-like extracellular matrix protein 2 (EFEMP2), mRNA. /FEA=mRNA /GEN=EFEMP2 /PROD=EGF-containing fibulin-like extracellular matrixprotein 2 /DB\_XREF=gi:8393298 /UG=Hs.6059 EGF-containing fibulin-like extracellular matrix protein 2 /FL=gb:AF093119.1 gb:AF109121.1 gb:NM\_016938.1

204379\_s\_at gb:NM\_000142.2 /DEF=Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) (FGFR3), transcript variant 1, mRNA. /FEA=mRNA /GEN=FGFR3 /PROD=fibroblast growth factor receptor 3, isoform 1precursor /DB\_XREF=gi:13112046 /UG=Hs.1420 fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) /FL=gb:NM\_000142.2 gb:M58051.1

213805\_at Consensus includes gb:AI692428 /FEA=EST /DB\_XREF=gi:4969768 /DB\_XREF=est:wd69c12.x1 /CLONE=IMAGE:2336854 /UG=Hs.184019 Homo sapiens clone 23551 mRNA sequence

207907\_at gb:NM\_003807.1 /DEF=Homo sapiens tumor necrosis factor (ligand) superfamily, member 14 (TNFSF14), mRNA. /FEA=mRNA /GEN=TNFSF14 /PROD=tumor necrosis factor (ligand) superfamily, member 14 /DB\_XREF=gi:4507600 /UG=Hs.129708 tumor necrosis factor (ligand) superfamily, member 14 /FL=gb:AF036581.1 gb:AF064090.1 gb:NM\_003807.1

204045\_at gb:NM\_004780.1 /DEF=Homo sapiens transcription elongation factor A (SII)-like 1 (TCEAL1), mRNA. /FEA=mRNA /GEN=TCEAL1 /PROD=transcription elongation factor A (SII)-like 1 /DB\_XREF=gi:4759215 /UG=Hs.95243 transcription elongation factor A (SII)-like 1 /FL=gb:BC000809.1 gb:NM\_004780.1 gb:M99701.1

# Affy Probe\_ID

## Sequence Description

210788\_s\_at gb:AF126782.1 /DEF=Homo sapiens retinal short-chain dehydrogenasereductase retSDR4 mRNA, complete cds. /FEA=mRNA /PROD=retinal short-chain dehydrogenasereductaseretSDR4 /DB\_XREF=gi:6318547 /UG=Hs.109201 CGI-86 protein /FL=gb:AF126782.1

204601\_at gb:NM\_014664.1 /DEF=Homo sapiens KIAA0615 gene product (KIAA0615), mRNA. /FEA=mRNA /GEN=KIAA0615 /PROD=KIAA0615 gene product /DB\_XREF=gi:7662203 /UG=Hs.323712 KIAA0615 gene product /FL=gb:AB014515.1 gb:NM\_014664.1

219179\_at gb:NM\_016651.2 /DEF=Homo sapiens heptacellular carcinoma novel gene-3 protein (LOC51339), mRNA. /FEA=mRNA /GEN=LOC51339 /PROD=heptacellular carcinoma novel gene-3 protein /DB\_XREF=gi:8393715 /UG=Hs.48950 heptacellular carcinoma novel gene-3 protein /FL=gb:AF251079.2 gb:NM\_016651.2

222241\_at Consensus includes gb:AK000001.1 /DEF=Homo sapiens mRNA for FLJ00001 protein, partial cds. /FEA=mRNA /GEN=FLJ00001 /PROD=FLJ00001 protein /DB\_XREF=gi:7209302 /UG=Hs.4914 Homo sapiens mRNA for FLJ00031 protein, partial cds

221210\_s\_at gb:NM\_030769.1 /DEF=Homo sapiens hypothetical protein similar to swine acylneuraminase lyase (C1ORF13), mRNA. /FEA=mRNA /GEN=C1ORF13 /PROD=hypothetical protein similar to swine acylneuraminase lyase /DB\_XREF=gi:13540532 /FL=gb:NM\_030769.1

215990\_s\_at Consensus includes gb:S67779.1 /DEF=BCL5=Cys2-His2 zinc-finger transcription factor human, liver, mRNA, 2600 nt. /FEA=mRNA /DB\_XREF=gi:459372 /UG=Hs.155024 B-cell CLL lymphoma 6 (zinc finger protein 51)

201650\_at gb:NM\_002276.1 /DEF=Homo sapiens keratin 19 (KRT19), mRNA. /FEA=mRNA /GEN=KRT19 /PROD=keratin 19 /DB\_XREF=gi:4504916 /UG=Hs.182265 keratin 19 /FL=gb:BC002539.1 gb:NM\_002276.1

208736\_at gb:AF004561.1 /DEF=Homo sapiens p21-Arc mRNA, complete cds. /FEA=mRNA /PROD=p21-Arc /DB\_XREF=gi:2209346 /UG=Hs.6895 actin related protein 23 complex, subunit 3 (21 kD) /FL=gb:AF004561.1 gb:AF006086.1 gb:NM\_005719.1

202598\_at gb:NM\_005979.1 /DEF=Homo sapiens S100 calcium-binding protein A13 (S100A13), mRNA. /FEA=mRNA /GEN=S100A13 /PROD=S100 calcium-binding protein A13 /DB\_XREF=gi:5174658 /UG=Hs.14331 S100 calcium-binding protein A13 /FL=gb:BC000632.1 gb:NM\_005979.1

203307\_at gb:NM\_005275.1 /DEF=Homo sapiens guanine nucleotide binding protein-like 1 (GNL1), mRNA. /FEA=mRNA /GEN=GNL1 /PROD=guanine nucleotide binding protein-like 1 /DB\_XREF=gi:4885288 /UG=Hs.83147 guanine nucleotide binding protein-like 1 /FL=gb:NM\_005275.1 gb:L25665.1

Affy Probe_ID	Sequence Description
205349_at	gb:NM_002068.1 /DEF=Homo sapiens guanine nucleotide binding protein (G protein), alpha 15 (Gq class) (GNA15), mRNA. /FEA=mRNA /GEN=GNA15 /PROD=guanine nucleotide binding protein (G protein), alpha 15 (Gq class) /DB_XREF=gi:4504038 /UG=Hs.73797 guanine nucleotide binding protein (G protein), alpha 15 (Gq class) /FL=gb:M63904.1 gb:NM_002068.1
201116_s_at	Consensus includes gb:AI922855 /FEA=EST /DB_XREF=gi:5658819 /DB_XREF=est:wo14h05.x1 /CLONE=IMAGE:2455353 /UG=Hs.75360 carboxypeptidase E /FL=gb:Nm_001873.1
200862_at	gb:NM_014762.1 /DEF=Homo sapiens seladin-1 (KIAA0018), mRNA. /FEA=mRNA /GEN=KIAA0018 /PROD=seladin-1 /DB_XREF=gi:13375617 /UG=Hs.75616 seladin-1 /FL=gb:AF261758.1 gb:BC004375.1 gb:NM_014762.1
218807_at	gb:NM_006113.2 /DEF=Homo sapiens vav 3 oncogene (VAV3), mRNA. /FEA=mRNA /GEN=VAV3 /PROD=vav 3 oncogene /DB_XREF=gi:7262390 /UG=Hs.267659 vav 3 oncogene /FL=gb:AF067817.1 gb:AF118887.1 gb:NM_006113.2
212514_x_at	Consensus includes gb:R60068 /FEA=EST /DB_XREF=gi:830763 /DB_XREF=est:yh12h07.s1 /CLONE=IMAGE:43138 /UG=Hs.147916 DEADH (Asp-Glu-Ala-AspHis) box polypeptide 3
208315_x_at	gb:NM_003300.1 /DEF=Homo sapiens TNF receptor-associated factor 3 (TRAF3), mRNA. /FEA=mRNA /GEN=TRAF3 /PROD=TNF receptor-associated factor 3 /DB_XREF=gi:4507678 /UG=Hs.297660 TNF receptor-associated factor 3 /FL=gb:Nm_003300.1 gb:U21092.1
201920_at	gb:NM_005415.2 /DEF=Homo sapiens solute carrier family 20 (phosphate transporter), member 1 (SLC20A1), mRNA. /FEA=mRNA /GEN=SLC20A1 /PROD=solute carrier family 20 (phosphate transporter), member 1 /DB_XREF=gi:7382462 /UG=Hs.78452 solute carrier family 20 (phosphate transporter), member 1 /FL=gb:L20859.1 gb:NM_005415.2
221974_at	Consensus includes gb:AW770748 /FEA=EST /DB_XREF=gi:7702795 /DB_XREF=est:hl90a07.x1 /CLONE=IMAGE:3009204 /UG=Hs.5022 imprinted in Prader-Willi syndrome
201516_at	gb:NM_003132.1 /DEF=Homo sapiens spermidine synthase (SRM), mRNA. /FEA=mRNA /GEN=SRM /PROD=spermidine synthase /DB_XREF=gi:4507208 /UG=Hs.76244 spermidine synthase /FL=gb:BC000309.1 gb:NM_003132.1 gb:M34338.1
209395_at	gb:M80927.1 /DEF=Human glycoprotein mRNA, complete cds. /FEA=mRNA /PROD=glycoprotein /DB_XREF=gi:348911 /UG=Hs.75184 chitinase 3-like 1 (cartilage glycoprotein-39) /FL=gb:M80927.1 gb:NM_001276.1



# Affy Probe\_ID

## Sequence Description

213616_at	Consensus includes gb:BF063896 /FEA=EST /DB_XREF=gi:10822806 /DB_XREF=est:7h90a05.x1 /CLONE=IMAGE:3323216 /UG=Hs.22981 DKFZP586M1523 protein
204521_at	gb:NM_013300.1 /DEF=Homo sapiens protein predicted by clone 23733 (HSU79274), mRNA. /FEA=mRNA /GEN=HSU79274 /PROD=protein predicted by clone 23733 /DB_XREF=gi:9558740 /UG=Hs.150555 protein predicted by clone 23733 /FL=gb:U79274.1 gb:NM_013300.1
213900_at	Consensus includes gb:AA524029 /FEA=EST /DB_XREF=gi:2264957 /DB_XREF=est:ng32f02.s1 /CLONE=IMAGE:936507 /UG=Hs.77889 Friedreich ataxia region gene X123
219682_s_at	gb:NM_016569.1 /DEF=Homo sapiens TBX3-iso protein (TBX3-iso), mRNA. /FEA=mRNA /GEN=TBX3-iso /PROD=TBX3-iso protein /DB_XREF=gi:7706728 /UG=Hs.332150 TBX3-iso protein /FL=gb:AF216750.1 gb:NM_016569.1
217513_at	Consensus includes gb:BG334495 /FEA=EST /DB_XREF=gi:13140933 /DB_XREF=est:602461128F1 /CLONE=IMAGE:4577718 /UG=Hs.228201 ESTs, Moderately similar to AF118094 25 PRO1992 H.sapiens
220432_s_at	gb:NM_016593.1 /DEF=Homo sapiens oxysterol 7alpha-hydroxylase (CYP39A1), mRNA. /FEA=mRNA /GEN=CYP39A1 /PROD=oxysterol 7alpha-hydroxylase /DB_XREF=gi:7706128 /UG=Hs.20766 oxysterol 7alpha-hydroxylase /FL=gb:AF237982.1 gb:NM_016593.1
202107_s_at	gb:NM_004526.1 /DEF=Homo sapiens minichromosome maintenance deficient (S. cerevisiae) 2 (mitotin) (MCM2), mRNA. /FEA=mRNA /GEN=MCM2 /PROD=minichromosome maintenance deficient (S. cerevisiae) 2 (mitotin) /DB_XREF=gi:4758705 /UG=Hs.57101 minichromosome maintenance deficient (S. cerevisiae) 2 (mitotin) /FL=gb:D83987.1 gb:NM_004526.1
204785_x_at	gb:NM_000874.1 /DEF=Homo sapiens interferon (alpha, beta and omega) receptor 2 (IFNAR2), mRNA. /FEA=mRNA /GEN=IFNAR2 /PROD=interferon (alpha, beta and omega) receptor 2 /DB_XREF=gi:4504600 /UG=Hs.86958 interferon (alpha, beta and omega) receptor 2 /FL=gb:NM_000874.1 gb:L41944.1
206134_at	gb:NM_014479.1 /DEF=Homo sapiens disintegrin protease (M12.219), mRNA. /FEA=mRNA /GEN=M12.219 /PROD=disintegrin protease /DB_XREF=gi:7657318 /UG=Hs.145296 disintegrin protease /FL=gb:NM_014479.1
213125_at	Consensus includes gb:AW007573 /FEA=EST /DB_XREF=gi:5856436 /DB_XREF=est:wt02e12.x1 /CLONE=IMAGE:2506318 /UG=Hs.43658 DKFZP586L151 protein
216836_s_at	Consensus includes gb:X03363.1 /DEF=Human c-erb-B-2 mRNA. /FEA=mRNA /DB_XREF=gi:31197 /UG=Hs.323910 v-erb-b2 avian erythroblastic leukemia viral oncogene homolog 2 (neuroglioblastoma derived oncogene homolog)

# Affy Probe\_ID

## Sequence Description

202796\_at gb:NM\_007286.1 /DEF=Homo sapiens synaptopodin (KIAA1029), mRNA. /FEA=mRNA /GEN=KIAA1029 /PROD=synaptopodin /DB\_XREF=gi:6005797 /UG=Hs.5307 synaptopodin /FL=gb:AB028952.1 gb:NM\_007286.1

219270\_at gb:NM\_024111.1 /DEF=Homo sapiens hypothetical protein MGC4504 (MGC4504), mRNA. /FEA=mRNA /GEN=MGC4504 /PROD=hypothetical protein MGC4504 /DB\_XREF=gi:13129125 /UG=Hs.155569 hypothetical protein MGC4504 /FL=gb:BC001847.1 gb:NM\_024111.1

216782\_at Consensus includes gb:AK026679.1 /DEF=Homo sapiens cDNA: FLJ23026 fis, clone LNG01738. /FEA=mRNA /DB\_XREF=gi:10439585 /UG=Hs.306863 Homo sapiens cDNA: FLJ23026 fis, clone LNG01738

219265\_at gb:NM\_024761.1 /DEF=Homo sapiens hypothetical protein FLJ13204 (FLJ13204), mRNA. /FEA=mRNA /GEN=FLJ13204 /PROD=hypothetical protein FLJ13204 /DB\_XREF=gi:13376100 /UG=Hs.128905 hypothetical protein FLJ13204 /FL=gb:NM\_024761.1

203726\_s\_at gb:NM\_000227.1 /DEF=Homo sapiens laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin) (LAMA3), mRNA. /FEA=mRNA /GEN=LAMA3 /PROD=laminin alpha 3 subunit precursor /DB\_XREF=gi:4557710 /UG=Hs.83450 laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin) /FL=gb:NM\_000227.1 gb:L34155.1

204824\_at gb:NM\_004435.1 /DEF=Homo sapiens endonuclease G (ENDOG), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=ENDOG /PROD=endonuclease G precursor /DB\_XREF=gi:4758269 /UG=Hs.182418 endonuclease G /FL=gb:BC004922.1 gb:NM\_004435.1

203049\_s\_at gb:NM\_014639.1 /DEF=Homo sapiens KIAA0372 gene product (KIAA0372), mRNA. /FEA=mRNA /GEN=KIAA0372 /PROD=KIAA0372 gene product /DB\_XREF=gi:7662077 /UG=Hs.170098 KIAA0372 gene product /FL=gb:AB002370.1 gb:NM\_014639.1

204641\_at gb:NM\_002497.1 /DEF=Homo sapiens NIMA (never in mitosis gene a)-related kinase 2 (NEK2), mRNA. /FEA=mRNA /GEN=NEK2 /PROD=NIMA (never in mitosis gene a)-related kinase 2 /DB\_XREF=gi:4505372 /UG=Hs.153704 NIMA (never in mitosis gene a)-related kinase 2 /FL=gb:NM\_002497.1 gb:U11050.1

219410\_at gb:NM\_018004.1 /DEF=Homo sapiens hypothetical protein FLJ10134 (FLJ10134), mRNA. /FEA=mRNA /GEN=FLJ10134 /PROD=hypothetical protein FLJ10134 /DB\_XREF=gi:8922242 /UG=Hs.104800 hypothetical protein FLJ10134 /FL=gb:NM\_018004.1

204417\_at gb:NM\_000153.1 /DEF=Homo sapiens galactosylceramidase (Krabbe disease) (GALC), mRNA. /FEA=mRNA /GEN=GALC /PROD=galactosylceramidase precursor /DB\_XREF=gi:4557612 /UG=Hs.273 galactosylceramidase (Krabbe disease) /FL=gb:L23116.1 gb:NM\_000153.1 gb:D25283.1

**Affy Probe\_ID**

**Sequence Description**

220168\_at gb:NM\_018272.1 /DEF=Homo sapiens hypothetical protein FLJ10921 (FLJ10921), mRNA. /FEA=mRNA /GEN=FLJ10921 /PROD=hypothetical protein FLJ10921 /DB\_XREF=gi:8922767 /UG=Hs.279012 hypothetical protein FLJ10921 /FL=gb:NM\_018272.1

209909\_s\_at gb:M19154.1 /DEF=Human transforming growth factor-beta-2 mRNA, complete cds. /FEA=mRNA /PROD=transforming growth factor beta 2 /DB\_XREF=gi:339549 /UG=Hs.169300 transforming growth factor, beta 2 /FL=gb:M19154.1

205805\_s\_at gb:NM\_005012.1 /DEF=Homo sapiens receptor tyrosine kinase-like orphan receptor 1 (ROR1), mRNA. /FEA=mRNA /GEN=ROR1 /PROD=receptor tyrosine kinase-like orphan receptor 1 /DB\_XREF=gi:4826867 /UG=Hs.274243 receptor tyrosine kinase-like orphan receptor 1 /FL=gb:M97675.1 gb:NM\_005012.1

214716\_at Consensus includes gb:AW504018 /FEA=EST /DB\_XREF=gi:7141685 /DB\_XREF=est:UI-HF-BN0-alh-f-03-0-UI.r1 /CLONE=IMAGE:3079804 /UG=Hs.20137 hypothetical protein DKFZp434P0116

214925\_s\_at Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA: FLJ22831 fis, clone KAIA4161. /FEA=mRNA /DB\_XREF=gi:10439356 /UG=Hs.321666 Homo sapiens cDNA: FLJ22831 fis, clone KAIA4161

205348\_s\_at gb:NM\_004411.1 /DEF=Homo sapiens dynein, cytoplasmic, intermediate polypeptide 1 (DNCCI), mRNA. /FEA=mRNA /GEN=DNCCI /PROD=dynein, cytoplasmic, intermediate polypeptide 1 /DB\_XREF=gi:4758177 /UG=Hs.65248 dynein, cytoplasmic, intermediate polypeptide 1 /FL=gb:AF063228.1 gb:NM\_004411.1

219109\_at gb:NM\_024532.1 /DEF=Homo sapiens hypothetical protein FLJ22724 (FLJ22724), mRNA. /FEA=mRNA /GEN=FLJ22724 /PROD=hypothetical protein FLJ22724 /DB\_XREF=gi:13375683 /UG=Hs.6783 hypothetical protein FLJ22724 /FL=gb:NM\_024532.1

203692\_s\_at Consensus includes gb:AI640363 /FEA=EST /DB\_XREF=gi:4703472 /DB\_XREF=est:wa17d03.x1 /CLONE=IMAGE:2298341 /UG=Hs.1189 E2F transcription factor 3 /FL=gb:NM\_001949.2

211628\_x\_at gb:J04755.1 /DEF=Human ferritin H processed pseudogene, complete cds. /FEA=mRNA /GEN=FTHP1 /DB\_XREF=gi:182512 /FL=gb:J04755.1

209768\_s\_at Consensus includes gb:AI860917 /FEA=EST /DB\_XREF=gi:5514533 /DB\_XREF=est:w156b07.x1 /CLONE=IMAGE:2428885 /UG=Hs.283743 glycoprotein Ib (platelet), beta polypeptide /FL=gb:L20860.1

211527\_x\_at gb:M27281.1 /DEF=Human vascular permeability factor mRNA, complete cds. /FEA=mRNA /DB\_XREF=gi:340300 /UG=Hs.73793 vascular endothelial growth factor /FL=gb:M27281.1

# Affy Probe ID

## Sequence Description

205241_at	gb:NM_005138.1 /DEF=Homo sapiens SCO (cytochrome oxidase deficient, yeast) homolog 2 (SCO2), nuclear gene encoding mitochondrial protein, mRNA. /FEA=mRNA /GEN=SCO2 /PROD=SCO (cytochrome oxidase deficient, yeast)homolog 2 /DB_XREF=gi:4826991 /UG=Hs.278431 SCO (cytochrome oxidase deficient, yeast) homolog 2 /FL=gb:NM_005138.1
219863_at	gb:NM_016323.1 /DEF=Homo sapiens cyclin-E binding protein 1 (LOC51191), mRNA. /FEA=mRNA /GEN=LOC51191 /PROD=cyclin-E binding protein 1 /DB_XREF=gi:7705930 /UG=Hs.26663 cyclin-E binding protein 1 /FL=gb:AB027289.1 gb:NM_016323.1
201091_s_at	Consensus includes gb:BE748755 /FEA=EST /DB_XREF=gi:10162747 /DB_XREF=est:601571933T1 /CLONE=IMAGE:3838737 /UG=Hs.278554 heterochromatin-like protein 1 /FL=gb:AF136630.1 gb:NM_016587.1
207433_at	gb:NM_000572.1 /DEF=Homo sapiens interleukin 10 (IL10), mRNA. /FEA=mRNA /GEN=IL10 /PROD=interleukin 10 /DB_XREF=gi:10835140 /UG=Hs.193717 interleukin 10 /FL=gb:NM_000572.1 gb:M57627.1
200622_x_at	Consensus includes gb:AV685208 /FEA=EST /DB_XREF=gi:10287071 /DB_XREF=est:AV685208 /CLONE=GKCCXA06 /UG=Hs.141011 calmodulin 3 (phosphorylase kinase, delta) /FL=gb:BC005137.1 gb:J04046.1 gb:NM_005184.1
203032_s_at	Consensus includes gb:AI363836 /FEA=EST /DB_XREF=gi:4123525 /DB_XREF=est:qy62g01.x1 /CLONE=IMAGE:2016624 /UG=Hs.75653 fumarate hydratase /FL=gb:BC003108.1 gb:M15502.1 gb:U59309.1 gb:U48857.1 gb:NM_000143.1
204237_at	gb:NM_016315.1 /DEF=Homo sapiens CED-6 protein (CED-6), mRNA. /FEA=mRNA /GEN=CED-6 /PROD=CED-6 protein /DB_XREF=gi:7705317 /UG=Hs.107056 CED-6 protein /FL=gb:AF200715.1 gb:AF191771.1 gb:NM_016315.1
216264_s_at	Consensus includes gb:X79683.1 /DEF=H.sapiens LAMB2 mRNA for beta2 laminin. /FEA=mRNA /GEN=LAMB2 /PROD=beta2S laminin chain /DB_XREF=gi:663206 /UG=Hs.90291 laminin, beta 2 (laminin S)
208710_s_at	Consensus includes gb:AI424923 /FEA=EST /DB_XREF=gi:4270841 /DB_XREF=est:tg19a07.x1 /CLONE=IMAGE:2109204 /UG=Hs.75056 adaptor-related protein complex 3, delta 1 subunit /FL=gb:AF002163.1
221541_at	gb:AL136861.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B044 (from clone DKFZp434B044); complete cds. /FEA=mRNA /GEN=DKFZp434B044 /PROD=hypothetical protein /DB_XREF=gi:12053226 /UG=Hs.262958 Homo sapiens mRNA; cDNA DKFZp434B044 (from clone DKFZp434B044); complete cds /FL=gb:AL136861.1

Affy Probe_ID	Sequence Description
206059_at	gb:NM_003430.1 /DEF=Homo sapiens zinc finger protein 91 (HPF7, HTF10) (ZNF91), mRNA. /FEA=mRNA /GEN=ZNF91 /PROD=zinc finger protein 91 (HPF7, HTF10) /DB_XREF=gi:4508040 /UG=Hs.8597 zinc finger protein 91 (HPF7, HTF10) /FL=gb:L11672.1 gb:NM_003430.1
200872_at	gb:NM_002966.1 /DEF=Homo sapiens S100 calcium-binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) (S100A10), mRNA. /FEA=mRNA /GEN=S100A10 /PROD=S100 calcium-binding protein A10 /DB_XREF=gi:4506760 /UG=Hs.119301 S100 calcium-binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) /FL=gb:M81457.1 gb:M38591.1 gb:NM_002966.1
203962_s_at	gb:NM_006393.1 /DEF=Homo sapiens nebulin (NEBL), mRNA. /FEA=mRNA /GEN=NEBL /PROD=nebulin /DB_XREF=gi:5453757 /UG=Hs.5025 nebulin /FL=gb:NM_006393.1
209617_s_at	gb:AF035302.1 /DEF=Homo sapiens clone 23717 mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:2661061 /UG=Hs.80220 catenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein) /FL=gb:NM_001332.1 gb:U72665.1 gb:AF035302.1 gb:U96136.1 gb:AB013805.1
207920_x_at	gb:NM_003410.1 /DEF=Homo sapiens zinc finger protein, X-linked (ZFX), mRNA. /FEA=mRNA /GEN=ZFX /PROD=zinc finger protein, X-linked /DB_XREF=gi:4507964 /UG=Hs.2074 zinc finger protein, X-linked /FL=gb:M30608.1 gb:NM_003410.1
202207_at	Consensus includes gb:BG435404 /FEA=EST /DB_XREF=gi:13341910 /DB_XREF=est:602507678F1 /CLONE=IMAGE:4605066 /UG=Hs.111554 ADP-ribosylation factor-like 7 /FL=gb:BC001051.1 gb:AB016811.1 gb:NM_005737.2
203853_s_at	gb:NM_012296.1 /DEF=Homo sapiens GRB2-associated binding protein 2 (GAB2), mRNA. /FEA=mRNA /GEN=GAB2 /PROD=GRB2-associated binding protein 2 /DB_XREF=gi:6912459 /UG=Hs.30687 GRB2-associated binding protein 2 /FL=gb:AB011143.1 gb:NM_012296.1
205759_s_at	gb:NM_004605.1 /DEF=Homo sapiens sulfotransferase family, cytosolic, 2B, member 1 (SULT2B1), mRNA. /FEA=mRNA /GEN=SULT2B1 /PROD=sulfotransferase family, cytosolic, 2B, member 1 /DB_XREF=gi:4759191 /UG=Hs.94581 sulfotransferase family, cytosolic, 2B, member 1 /FL=gb:U92314.1 gb:NM_004605.1
221811_at	Consensus includes gb:BF033007 /FEA=EST /DB_XREF=gi:10740719 /DB_XREF=est:601455980F1 /CLONE=IMAGE:3859819 /UG=Hs.91668 Homo sapiens clone PP1498 unknown mRNA

**Affy Probe ID**

**Sequence Description**

200704_at	gb:AB034747.1 /DEF=Homo sapiens SIMPLE mRNA for small integral membrane protein of lysosomal endosome, complete cds. /FEA=mRNA /GEN=SIMPLE /PROD=small integral membrane protein of lysosomal endosome /DB_XREF=gi:12862475 /UG=Hs.76507 LPS-induced TNF-alpha factor /FL=gb:AB034747.1 gb:U77396.1 gb:AF010312.1 gb:NM_004862.1
203484_at	gb:NM_014302.1 /DEF=Homo sapiens Sec61 gamma (SEC61G), mRNA. /FEA=mRNA /GEN=SEC61G /PROD=Sec61 gamma /DB_XREF=gi:7657545 /UG=Hs.9950 Sec61 gamma /FL=gb:AF054184.1 gb:NM_014302.1
200021_at	gb:NM_005507.1 /DEF=Homo sapiens cofilin 1 (non-muscle) (CFL1), mRNA. /FEA=mRNA /GEN=CFL1 /PROD=cofilin 1 (non-muscle) /DB_XREF=gi:5031634 /UG=Hs.180370 cofilin 1 (non-muscle) /FL=gb:NM_005507.1
218901_at	gb:NM_020353.1 /DEF=Homo sapiens phospholipid scramblase 4 (LOC57088), mRNA. /FEA=mRNA /GEN=LOC57088 /PROD=phospholipid scramblase 4 /DB_XREF=gi:9966818 /UG=Hs.182538 phospholipid scramblase 4 /FL=gb:AF199023.1 gb:NM_020353.1
207619_at	gb:NM_001525.1 /DEF=Homo sapiens hypocretin (orexin) receptor 1 (HCRTR1), mRNA. /FEA=mRNA /GEN=HCRTR1 /PROD=orexin receptor 1 /DB_XREF=gi:4557636 /UG=Hs.150968 hypocretin (orexin) receptor 1 /FL=gb:AF041243.1 gb:NM_001525.1
206157_at	gb:NM_002852.1 /DEF=Homo sapiens pentaxin-related gene, rapidly induced by IL-1 beta (PTX3), mRNA. /FEA=mRNA /GEN=PTX3 /PROD=pentaxin-related gene, rapidly induced by IL-1 beta /DB_XREF=gi:4506332 /UG=Hs.2050 pentaxin-related gene, rapidly induced by IL-1 beta /FL=gb:M31166.1 gb:NM_002852.1
217902_s_at	gb:NM_004667.2 /DEF=Homo sapiens hect domain and RLD 2 (HERC2), mRNA. /FEA=mRNA /GEN=HERC2 /PROD=hect domain and RLD 2 /DB_XREF=gi:5729867 /UG=Hs.266933 hect domain and RLD 2 /FL=gb:AF071172.3 gb:NM_004667.2
200729_s_at	gb:NM_005722.1 /DEF=Homo sapiens ARP2 (actin-related protein 2, yeast) homolog (ACTR2), mRNA. /FEA=mRNA /GEN=ACTR2 /PROD=ARP2 (actin-related protein 2, yeast) homolog /DB_XREF=gi:5031570 /UG=Hs.42915 ARP2 (actin-related protein 2, yeast) homolog /FL=gb:AF006082.1 gb:NM_005722.1
213547_at	Consensus includes gb:AB014567.1 /DEF=Homo sapiens mRNA for KIAA0667 protein, partial cds. /FEA=mRNA /GEN=KIAA0667 /PROD=KIAA0667 protein /DB_XREF=gi:3327147 /UG=Hs.154740 TBP-interacting protein - Consensus includes gb:AA88589 /FEA=EST /DB_XREF=gi:3004264 /DB_XREF=est:oe68e10.s1
209425_at	/CLONE=IMAGE:1416810 /UG=Hs.128749 alpha-methylacyl-CoA racemase /FL=gb:AF047020.1 gb:AF158378.1 gb:NM_014324.1

Affy Probe_ID	Sequence Description
205525_at	gb:NM_018495.3 /DEF=Homo sapiens NAG22 protein (LOC55873), mRNA. /FEA=mRNA /GEN=LOC55873 /PROD=NAG22 protein /DB_XREF=gi:13236500 /UG=Hs.283080 NAG22 protein /FL=gb:AF247820.3 gb:NM_018495.3
219508_at	gb:NM_004751.1 /DEF=Homo sapiens glucosaminyl (N-acetyl) transferase 3, mucin type (GCNT3), mRNA. /FEA=mRNA /GEN=GCNT3 /PROD=glucosaminyl (N-acetyl) transferase 3, mucintype /DB_XREF=gi:4758421 /UG=Hs.194710 glucosaminyl (N-acetyl) transferase 3, mucin type /FL=gb:AF102542.1 gb:AF038650.1 gb:NM_004751.1
218916_at	gb:NM_024671.1 /DEF=Homo sapiens hypothetical protein FLJ23436 (FLJ23436), mRNA. /FEA=mRNA /GEN=FLJ23436 /PROD=hypothetical protein FLJ23436 /DB_XREF=gi:13375931 /UG=Hs.85658 hypothetical protein FLJ23436 /FL=gb:NM_024671.1
219811_at	gb:NM_022720.1 /DEF=Homo sapiens DiGeorge syndrome critical region gene 8 (DGCR8), mRNA. /FEA=mRNA /GEN=DGCR8 /PROD=DiGeorge syndrome critical region gene 8 /DB_XREF=gi:12225241 /UG=Hs.301855 DiGeorge syndrome critical region gene 8 /FL=gb:NM_022720.1 gb:AF165527.1
215129_at	Consensus includes gb:AJ000008.1 /DEF=Homo sapiens mRNA for C2 domain containing PI3-kinase. /FEA=mRNA /PROD=PI3-kinase /DB_XREF=gi:3288456 /UG=Hs.22500 phosphoinositide-3-kinase, class 2, gamma polypeptide
215169_at	Consensus includes gb:BE885244 /FEA=EST /DB_XREF=gi:10334020 /DB_XREF=est:60150631 /CLONE=IMAGE:3907947 /UG=Hs.214646 KIAA0447 gene product
212723_at	Consensus includes gb:AK021780.1 /DEF=Homo sapiens cDNA FLJ11718 fis, clone HEMBA1005252, highly similar to Homo sapiens mRNA for KIAA0585 protein. /FEA=mRNA /DB_XREF=gi:10433034 /UG=Hs.726660 phosphatidylserine receptor
210495_x_at	gb:AF130095.1 /DEF=Homo sapiens clone FLC0562 PRO2841 mRNA, complete cds. /FEA=mRNA /PROD=PRO2841 /DB_XREF=gi:11493493 /UG=Hs.287820 fibronectin 1 /FL=gb:AF130095.1
210916_s_at	gb:AF098641.1 /DEF=Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds. /FEA=mRNA /GEN=CD44 /PROD=CD44 isoform RC /DB_XREF=gi:3832517 /UG=Hs.306278 Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds /FL=gb:AF098641.1
220321_s_at	gb:NM_024584.1 /DEF=Homo sapiens hypothetical protein FLJ13646 (FLJ13646), mRNA. /FEA=mRNA /GEN=FLJ13646 /PROD=hypothetical protein FLJ13646 /DB_XREF=gi:13375767 /UG=Hs.21081 hypothetical protein FLJ13646 /FL=gb:NM_024584.1

# **Affy Probe\_ID**

## **Sequence Description**

207224\_s\_at gb:NM\_016543.1 /DEF=Homo sapiens D-siglec precursor, (LOC51766), mRNA. /FEA=mRNA /GEN=LOC51766 /PROD=D-siglec precursor, /DB\_XREF=gi:7706570 /UG=Hs.149250 D-siglec precursor, /FL=gb:AF178981.1 gb:NM\_016543.1

218025\_s\_at gb:NM\_006117.1 /DEF=Homo sapiens peroxisomal D3,D2-enoyl-CoA isomerase (PECI), mRNA. /FEA=mRNA /GEN=PECI /PROD=peroxisomal D3,D2-enoyl-CoA isomerase /DB\_XREF=gi:5174624 /UG=Hs.15250 peroxisomal D3,D2-enoyl-CoA isomerase /FL=gb:AL136642.1 gb:BC002668.1 gb:AF069301.1 gb:AF153612.1 gb:NM\_006117.1 gb:AF244138.1

212067\_s\_at Consensus includes gb:AL573058 /FEA=EST /DB\_XREF=gi:12931931 /DB\_XREF=est:AL573058 /CLONE=CS0DI014YC05 (3 prime) /UG=Hs.1279 complement component 1, r subcomponent

200887\_s\_at gb:NM\_007315.1 /DEF=Homo sapiens signal transducer and activator of transcription 1, 91kD (STAT1), mRNA. /FEA=mRNA /GEN=STAT1 /PROD=signal transducer and activator of transcription1, 91kD /DB\_XREF=gi:6274551 /UG=Hs.21486 signal transducer and activator of transcription 1, 91kD /FL=gb:M97935.1 gb:NM\_007315.1

214996\_at Consensus includes gb:AL079294.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362780. /FEA=mRNA /DB\_XREF=gi:5102751 /UG=Hs.29952 Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362780

211596\_s\_at gb:AB050468.1 /DEF=Homo sapiens mRNA for membrane glycoprotein LIG-1, complete cds. /FEA=mRNA /GEN=lig-1 /PROD=membrane glycoprotein LIG-1 /DB\_XREF=gi:13537354 /FL=gb:AB050468.1

214967\_at Consensus includes gb:AU146983 /FEA=EST /DB\_XREF=gi:11008504 /DB\_XREF=est:AU146983 /CLONE=HEMBB1002049 /UG=Hs.262212 Homo sapiens cDNA FLJ12055 fis, clone HEMBB1002049

202874\_s\_at gb:NM\_001695.1 /DEF=Homo sapiens ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD (ATP6C), mRNA. /FEA=mRNA /GEN=ATP6C /PROD=ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /DB\_XREF=gi:4502314 /UG=Hs.86905 ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD /FL=gb:NM\_001695.1

43427\_at Cluster Incl. AI970898:wr21c03.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2488324 /clone\_end=3 /gb=AI970898 /gi=5767724 /ug=Hs.234898 /len=382

211862\_x\_at gb:AF015451.1 /DEF=Homo sapiens Usurpin-beta mRNA, complete cds. /FEA=CDS /PROD=Usurpin-beta /DB\_XREF=gi:3133282 /UG=Hs.195175 CASP8 and FADD-like apoptosis regulator /FL=gb:AF015451.1

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NO.: DERM1120



**Affy Probe ID**

**Sequence Description**

202727_s_at	gb:NM_000416.1 /DEF=Homo sapiens interferon gamma receptor 1 (IFNGR1), mRNA. /FEA=mRNA /GEN=IFNGR1 /PROD=interferon gamma receptor 1 /DB_XREF=gi:4557879 /UG=Hs.180866 interferon gamma receptor 1 /FL=gb:BC005333.1 gb:J03143.1 gb:NM_000416.1
204563_at	gb:NM_000655.2 /DEF=Homo sapiens selectin L (lymphocyte adhesion molecule 1) (SELL), mRNA. /FEA=mRNA /GEN=SELL /PROD=selectin L /DB_XREF=gi:5713320 /UG=Hs.82848 selectin L (lymphocyte adhesion molecule 1) /FL=gb:M25280.1 gb:NM_000655.2
200733_s_at	gb:U48296.1 /DEF=Homo sapiens protein tyrosine phosphatase PTPCAAX1 (hPTPCAAX1) mRNA, complete cds. /FEA=mRNA /GEN=hPTPCAAX1 /PROD=protein tyrosine phosphatase PTPCAAX1 /DB_XREF=gi:1777754 /UG=Hs.227777 protein tyrosine phosphatase type IVA, member 1 /FL=gb:U48296.1 gb:NM_003463.1 Consensus includes gb:AI924903 /FEA=EST /DB_XREF=gi:5660867 /DB_XREF=est:wn25e04.x1 /CLONE=IMAGE:2446494 /UG=Hs.151706 KIAA0134 gene product /FL=gb:D50924.1 gb:NM_014681.1 Consensus includes gb:W68084 /FEA=EST /DB_XREF=gi:1376954 /DB_XREF=est:zd42f12.s1 /CLONE=IMAGE:343343 /UG=Hs.5599 EGF-like-domain, multiple 5
204815_s_at	
212830_at	gb:NM_017938.1 /DEF=Homo sapiens hypothetical protein FLJ20716 (FLJ20716), mRNA. /FEA=mRNA /GEN=FLJ20716 /PROD=hypothetical protein FLJ20716 /DB_XREF=gi:8923642 /UG=Hs.144517 hypothetical protein FLJ20716 /FL=gb:NM_017938.1
212070_at	Consensus includes gb:AL554008 /FEA=EST /DB_XREF=gi:12894377 /DB_XREF=est:AL554008 /CLONE=CS0DI079YN13 (5 prime) /UG=Hs.6527 G protein-coupled receptor 56
219013_at	gb:NM_022087.1 /DEF=Homo sapiens hypothetical protein FLJ21634 (FLJ21634), mRNA. /FEA=mRNA /GEN=FLJ21634 /PROD=hypothetical protein FLJ21634 /DB_XREF=gi:11545800 /UG=Hs.97056 hypothetical protein FLJ21634 /FL=gb:NM_022087.1
210569_s_at	gb:AF247180.1 /DEF=Homo sapiens sialic acid-binding immunoglobulin-like lectin-9 mRNA, complete cds. /FEA=mRNA /PROD=sialic acid-binding immunoglobulin-like lectin-9 /DB_XREF=gi:9392623 /UG=Hs.245828 sialic acid binding Ig-like lectin 9 /FL=gb:AF227924.1 gb:AF247180.1
202371_at	gb:NM_024863.1 /DEF=Homo sapiens hypothetical protein FLJ21174 (FLJ21174), mRNA. /FEA=mRNA /GEN=FLJ21174 /PROD=hypothetical protein FLJ21174 /DB_XREF=gi:13376293 /UG=Hs.194329 hypothetical protein FLJ21174 /FL=gb:AF271783.1 gb:NM_024863.1

**Affy Probe ID**

**Sequence Description**

207198\_s\_at gb:NM\_004987.2 /DEF=Homo sapiens LIM and senescent cell antigen-like domains 1 (LIMS1), mRNA. /FEA=mRNA /GEN=LIMS1 /PROD=LIM and senescent cell antigen-like domains 1 /DB\_XREF=gi:13518025 /UG=Hs.112378 LIM and senescent cell antigen-like domains 1 /FL=gb:U09284.2 gb:NM\_004987.2

209301\_at gb:M36532.1 /DEF=Human carbonic anhydrase II mRNA, complete cds. /FEA=mRNA /GEN=CA2 /DB\_XREF=gi:179794 /UG=Hs.155097 carbonic anhydrase II /FL=gb:J03037.1 gb:M36532.1 gb:NM\_000067.1

201522\_x\_at gb:NM\_003097.2 /DEF=Homo sapiens small nuclear ribonucleoprotein polypeptide N (SNRPN), transcript variant 1, mRNA. /FEA=mRNA /GEN=SNRPN /PROD=small nuclear ribonucleoprotein polypeptide N /DB\_XREF=gi:13027651 /UG=Hs.48375 small nuclear ribonucleoprotein polypeptide N /FL=gb:U41303.1

202404\_s\_at gb:NM\_003097.2 gb:BC003180.1 gb:J04615.1

202404\_s\_at gb:NM\_000089.1 /DEF=Homo sapiens collagen, type I, alpha 2 (COL1A2), mRNA. /FEA=mRNA /GEN=COL1A2 /PROD=collagen, type I, alpha 2 /DB\_XREF=gi:4502946 /UG=Hs.179573 collagen, type I, alpha 2 /FL=gb:J03464.1

202404\_s\_at gb:NM\_000089.1

211474\_s\_at gb:BC004948.1 /DEF=Homo sapiens, clone MGC:10846, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10846) /DB\_XREF=gi:13436313 /UG=Hs.41072 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 /FL=gb:BC004948.1

206045\_s\_at gb:NM\_003787.1 /DEF=Homo sapiens nucleolar protein 4 (NOL4), mRNA. /FEA=mRNA /GEN=NOL4 /PROD=nucleolar protein 4 /DB\_XREF=gi:4505420 /UG=Hs.6414 nucleolar protein 4 /FL=gb:AB017800.1

206045\_s\_at gb:NM\_003787.1

214978\_s\_at Consensus includes gb:AK023365.1 /DEF=Homo sapiens cDNA FLJ13303 fis, clone OVARC1001372, highly similar to Homo sapiens liprin-alpha4 mRNA. /FEA=mRNA /DB\_XREF=gi:10435269 /UG=Hs.300704 Homo sapiens cDNA FLJ13303 fis, clone OVARC1001372, highly similar to Homo sapiens liprin-alpha4 mRNA

210029\_at gb:M34455.1 /DEF=Human interferon-gamma-inducible indoleamine 2,3-dioxygenase (IDO) mRNA, complete cds. /FEA=mRNA /DB\_XREF=gi:185790 /UG=Hs.840 indoleamine-pyrrole 2,3 dioxygenase /FL=gb:M34455.1

210029\_at gb:NM\_002164.1

200971\_s\_at gb:NM\_014445.1 /DEF=Homo sapiens stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4 (SERP1), mRNA. /FEA=mRNA /GEN=SERP1 /PROD=stress-associated endoplasmic reticulum protein1; ribosome associated membrane protein 4 /DB\_XREF=gi:7657551 /UG=Hs.76698 stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4 /FL=gb:AL136807.1 gb:AF136975.1

200971\_s\_at gb:AB022427.1 gb:NM\_014445.1

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## Sequence Description

213713\_s\_at Consensus includes gb:R48779 /FEA=EST /DB\_XREF=gi:810805 /DB\_XREF=est:yj69c07.s1 /CLONE=IMAGE:153996 /UG=Hs.293845 ESTs, Weakly similar to B32688 beta-galactosidase-related protein H.sapiens

212946\_at Consensus includes gb:AK025432.1 /DEF=Homo sapiens cDNA: FLJ21779 fis, clone HEP00210. /FEA=mRNA /DB\_XREF=gi:10437943 /UG=Hs.151385 KIAA0564 protein

203162\_s\_at gb:NM\_005886.1 /DEF=Homo sapiens katanin p80 (WD40-containing) subunit B 1 (KATNB1), mRNA. /FEA=mRNA /GEN=KATNB1 /PROD=katanin p80 subunit B 1 /DB\_XREF=gi:5031816 /UG=Hs.275675 katanin p80 (WD40-containing) subunit B 1 /FL=gb:BC001353.1 gb:AF052432.1 gb:NM\_005886.1

202064\_s\_at gb:AF052059.1 /DEF=Homo sapiens SEL1L (SEL1L) mRNA, complete cds. /FEA=mRNA /GEN=SEL1L /PROD=SEL1L /DB\_XREF=gi:6851088 /UG=Hs.181300 sel-1 (suppressor of lin-12, C.elegans)-like /FL=gb:NM\_005065.1 gb:AB020335.1 gb:AF052059.1 gb:U11037.1

219735\_s\_at gb:NM\_014553.1 /DEF=Homo sapiens LBP protein (LBP-9), mRNA. /FEA=mRNA /GEN=LBP-9 /PROD=LBP protein /DB\_XREF=gi:7657298 /UG=Hs.114747 LBP protein; likely ortholog of mouse CRTR-1 /FL=gb:AF198488.1 gb:NM\_014553.1

212008\_at Consensus includes gb:N29889 /FEA=EST /DB\_XREF=gi:1148409 /DB\_XREF=est:yy11e11.s1 /CLONE=IMAGE:270956 /UG=Hs.77495 UBX domain-containing 1.

221795\_at Consensus includes gb:AI346341 /FEA=EST /DB\_XREF=gi:4083547 /DB\_XREF=est:qp50b09.x1 /CLONE=IMAGE:1926425 /UG=Hs.298998 Homo sapiens, Similar to hypothetical protein FLJ20093, clone MGC:1076, mRNA, complete cds

209821\_at gb:AB024518.1 /DEF=Homo sapiens mRNA for DVS27-related protein, complete cds. /FEA=mRNA /GEN=DVS27 /PROD=DVS27-related protein /DB\_XREF=gi:4520327 /UG=Hs.58589 glycogenin 2 /FL=gb:AB024518.1

205602\_x\_at gb:NM\_002783.1 /DEF=Homo sapiens pregnancy specific beta-1-glycoprotein 7 (PSG7), mRNA. /FEA=mRNA /GEN=PSG7 /PROD=pregnancy specific beta-1-glycoprotein 7 /DB\_XREF=gi:4506176 /UG=Hs.225932 pregnancy specific beta-1-glycoprotein 7 /FL=gb:NM\_002783.1 gb:U18467.1

211144\_x\_at gb:M30894.1 /DEF=Human T-cell receptor Ti rearranged gamma-chain mRNA V-J-C region, complete cds. /FEA=mRNA /GEN=CD3G /DB\_XREF=gi:339406 /UG=Hs.112259 T cell receptor gamma locus /FL=gb:M30894.1

215231\_at Consensus includes gb:AU144309 /FEA=EST /DB\_XREF=gi:11005830 /DB\_XREF=est:AU144309 /CLONE=HEMBA1001533 /UG=Hs.55830 Homo sapiens cDNA FLJ10068 fis, clone HEMBA1001533

# Affy Probe\_ID

## Sequence Description

203245\_s\_at gb:BC002791.1 /DEF=Homo sapiens, homologous to yeast nitrogen permease (candidate tumor suppressor), clone MGC:3527, mRNA, complete cds. /FEA=mRNA /PROD=homologous to yeast nitrogen permease (candidate tumor suppressor) /DB\_XREF=gi:12803888 /UG=Hs.169780 homologous to yeast nitrogen permease (candidate tumor suppressor) /FL=gb:BC002791.1 gb:AF040707.1 gb:Nm\_006545.1

205695\_at gb:Nm\_006843.1 /DEF=Homo sapiens serine dehydratase (SDS), mRNA. /FEA=mRNA /GEN=SDS /PROD=serine dehydratase /DB\_XREF=gi:5803160 /UG=Hs.76751 serine dehydratase /FL=gb:J05037.1 gb:Nm\_006843.1

221928\_at Consensus includes gb:AI057637 /FEA=EST /DB\_XREF=gi:3331503 /DB\_XREF=est:oy31h06.x1 /CLONE=IMAGE:1667483 /UG=Hs.234898 ESTs, Weakly similar to 2109260A B cell growth factor H.sapiens

211191\_at gb:AF054818.1 /DEF=Homo sapiens leukocyte differentiation antigen CD84 isoform CD84s (CD84) mRNA, complete cds. /FEA=mRNA /GEN=CD84 /PROD=leukocyte differentiation antigen CD84 isoform CD84s

201141\_at /DB\_XREF=gi:6650111 /UG=Hs.137548 CD84 antigen (leukocyte antigen) /FL=gb:AF054818.1

gb:Nm\_002510.1 /DEF=Homo sapiens glycoprotein (transmembrane) nmb (GPNMB), mRNA. /FEA=mRNA /GEN=GPNMB /PROD=glycoprotein (transmembrane) nmb /DB\_XREF=gi:4505404 /UG=Hs.82226 glycoprotein (transmembrane) nmb /FL=gb:AF322909.1 gb:Nm\_002510.1

217299\_s\_at Consensus includes gb:AK001017.1 /DEF=Homo sapiens cDNA FLJ10155 fis, clone HEMBA1003433, highly similar to Homo sapiens gene for NBS1. /FEA=mRNA /DB\_XREF=gi:7022031 /UG=Hs.25812 Nijmegen breakage syndrome 1 (nibrin)

213338\_at Consensus includes gb:BF062629 /FEA=EST /DB\_XREF=gi:10821539 /DB\_XREF=est:7h62h07.x1 /CLONE=IMAGE:3320605 /UG=Hs.35861 DKFZP586E1621 protein

213416\_at Consensus includes gb:BG532690 /FEA=EST /DB\_XREF=gi:13524229 /DB\_XREF=est:602580263F1

220349\_s\_at /CLONE=IMAGE:4718390 /UG=Hs.40034 integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor) gb:Nm\_022759.1 /DEF=Homo sapiens hypothetical protein FLJ21865 (FLJ21865), mRNA. /FEA=mRNA /GEN=FLJ21865 /PROD=hypothetical protein FLJ21865 /DB\_XREF=gi:12232428 /UG=Hs.29288 hypothetical protein FLJ21865 /FL=gb:Nm\_022759.1

219807\_x\_at gb:Nm\_016154.1 /DEF=Homo sapiens ras-related GTP-binding protein 4b (RAB4B), mRNA. /FEA=mRNA /GEN=RAB4B /PROD=ras-related GTP-binding protein 4b /DB\_XREF=gi:7706672 /UG=Hs.279771 Homo sapiens TR00071289\_m (RAB4B), mRNA /FL=gb:AF165522.1 gb:Nm\_016154.1

212566\_at Consensus includes gb:AL523310 /FEA=EST /DB\_XREF=gi:12786803 /DB\_XREF=est:AL523310 /CLONE=CS0DC001YN06 (3 prime) /UG=Hs.150580 putative translation initiation factor

Affy Probe_ID	Sequence Description
212170_at	Consensus includes gb:BF447705 /FEA=EST /DB_XREF=gi:11512843 /DB_XREF=est:7q94d07.x1 /CLONE=IMAGE:3706021 /UG=Hs.180895 putative brain nuclearly-targeted protein
220007_at	gb:Nm_024770.1 /DEF=Homo sapiens hypothetical protein FLJ13984 (FLJ13984), mRNA. /FEA=mRNA /GEN=FLJ13984 /PROD=hypothetical protein FLJ13984 /DB_XREF=gi:13376116 /UG=Hs.135146 hypothetical protein FLJ13984 /FL=gb:Nm_024770.1
201489_at	gb:BC005020.1 /DEF=Homo sapiens, peptidylprolyl isomerase F (cyclophilin F), clone MGC:11022, mRNA, complete cds. /FEA=mRNA /PROD=peptidylprolyl isomerase F (cyclophilin F) /DB_XREF=gi:13477126 /UG=Hs.173125 peptidylprolyl isomerase F (cyclophilin F) /FL=gb:BC005020.1 gb:M80254.1 gb:Nm_005729.1
218820_at	gb:Nm_020215.1 /DEF=Homo sapiens hypothetical protein DKFZp761F2014 (DKFZp761F2014), mRNA. /FEA=mRNA /GEN=DKFZp761F2014 /PROD=hypothetical protein DKFZp761F2014 /DB_XREF=gi:9910205 /UG=Hs.6434 hypothetical protein DKFZp761F2014 /FL=gb:Nm_020215.1
212310_at	Consensus includes gb:D87742.1 /DEF=Human mRNA for KIAA0268 gene, partial cds. /FEA=mRNA /GEN=KIAA0268 /DB_XREF=gi:1665824 /UG=Hs.241552 KIAA0268 protein
216274_s_at	Consensus includes gb:N99438 /FEA=EST /DB_XREF=gi:1270864 /DB_XREF=est:yz96c06.r1 /CLONE=IMAGE:290890 /UG=Hs.9534 signal peptidase complex (18kD)
201118_at	gb:Nm_002631.1 /DEF=Homo sapiens phosphogluconate dehydrogenase (PGD), mRNA. /FEA=mRNA /GEN=PGD /PROD=phosphogluconate dehydrogenase /DB_XREF=gi:4505758 /UG=Hs.75888 phosphogluconate dehydrogenase /FL=gb:BC000368.1 gb:Nm_002631.1 gb:U30255.1
219885_at	gb:Nm_018042.1 /DEF=Homo sapiens hypothetical protein FLJ10260 (FLJ10260), mRNA. /FEA=mRNA /GEN=FLJ10260 /PROD=hypothetical protein FLJ10260 /DB_XREF=gi:8922317 /UG=Hs.105859 hypothetical protein FLJ10260 /FL=gb:Nm_018042.1
217597_x_at	Consensus includes gb:AI344141 /FEA=EST /DB_XREF=gi:4081347 /DB_XREF=est:tc02d09.x1 /CLONE=IMAGE:2062673 /UG=Hs.229850 ESTs, Highly similar to G01251 Rat protein H.sapiens
202074_s_at	gb:Nm_021980.1 /DEF=Homo sapiens tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein (FIP2), mRNA. /FEA=mRNA /GEN=FIP2 /PROD=tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtininteracting protein L; transcription factor IIIA-interacting protein /DB_XREF=gi:11415041 /UG=Hs.278898 tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein /FL=gb:Nm_021980.1

# Affy Probe\_ID

## Sequence Description

209598\_at gb:AB020690.1 /DEF=Homo sapiens mRNA for KIAA0883 protein, complete cds. /FEA=mRNA /GEN=KIAA0883 /PROD=KIAA0883 protein /DB\_XREF=gi:4240254 /UG=Hs.7782 paraneoplastic antigen MA2 /FL=gb:AF286487.1 gb:NM\_007257.1 gb:AB020690.1

203549\_s\_at

gb:NM\_000237.1 /DEF=Homo sapiens lipoprotein lipase (LPL), mRNA. /FEA=mRNA /GEN=LPL /PROD=lipoprotein lipase precursor /DB\_XREF=gi:4557726 /UG=Hs.180878 lipoprotein lipase /FL=gb:M15856.1 gb:NM\_000237.1 Consensus includes gb:AU158490 /FEA=EST /DB\_XREF=gi:11020011 /DB\_XREF=est:AU158490 /CLONE=PLACE300042 /UG=Hs.303632 Human DNA sequence from clone RP11-110H4 on chromosome 5 Contains a pseudogene similar to GUSB (glucuronidase, beta), a gene for a membrane protein, ESTs, STSs, GSSs and CpG islands

209166\_s\_at

gb:U68567.1 /DEF=Human lysosomal acid alpha-mannosidase mRNA, complete cds. /FEA=mRNA /PROD=lysosomal acid alpha-mannosidase /DB\_XREF=gi:1658373 /UG=Hs.279854 mannosidase, alpha, class 2B, member 1 /FL=gb:NM\_000528.1 gb:BC000736.1 gb:U05572.1 gb:U68567.1 gb:U60266.1

203043\_at

gb:NM\_004729.1 /DEF=Homo sapiens Ac-like transposable element (ALTE), mRNA. /FEA=mRNA /GEN=ALTE /PROD=Ac-like transposable element /DB\_XREF=gi:4759257 /UG=Hs.9933 Ac-like transposable element /FL=gb:AB018328.1 gb:NM\_004729.1

217914\_at

gb:NM\_017901.1 /DEF=Homo sapiens two-pore channel 1, homolog (KIAA1169), mRNA. /FEA=mRNA /GEN=KIAA1169 /PROD=two-pore channel 1, homolog /DB\_XREF=gi:8923566 /UG=Hs.26440 two-pore channel 1, homolog /FL=gb:NM\_017901.1

204613\_at

gb:NM\_002661.1 /DEF=Homo sapiens phospholipase C, gamma 2 (phosphatidylinositol-specific) (PLCG2), mRNA. /FEA=mRNA /GEN=PLCG2 /PROD=phospholipase C, gamma 2 (phosphatidylinositol-specific) /DB\_XREF=gi:4505870 /UG=Hs.75648 phospholipase C, gamma 2 (phosphatidylinositol-specific) /FL=gb:M37238.1 gb:NM\_002661.1

205866\_at

gb:NM\_003665.1 /DEF=Homo sapiens ficolin (collagenfibrinogen domain-containing) 3 (Hakata antigen) (FCN3), mRNA. /FEA=mRNA /GEN=FCN3 /PROD=ficolin 3 precursor /DB\_XREF=gi:4504330 /UG=Hs.21415 ficolin (collagenfibrinogen domain-containing) 3 (Hakata antigen) /FL=gb:D88587.1 gb:NM\_003665.1

221295\_at

gb:NM\_001279.1 /DEF=Homo sapiens cell death-inducing DFFA-like effector a (CIDEA), mRNA. /FEA=CDS /GEN=CIDEA /PROD=cell death-inducing DFFA-like effector a /DB\_XREF=gi:4557464 /UG=Hs.249129 cell death-inducing DFFA-like effector a /FL=gb:AF041378.1 gb:NM\_001279.1

# Affy Probe\_ID

## Sequence Description

212814\_at Consensus includes gb:AB020635.1 /DEF=Homo sapiens mRNA for KIAA0828 protein, partial cds. /FEA=mRNA /GEN=KIAA0828 /PROD=KIAA0828 protein /DB\_XREF=gi:4240144 /UG=Hs.4984 KIAA0828 protein

219563\_at gb:NM\_024633.1 /DEF=Homo sapiens hypothetical protein FLJ21276 (FLJ21276), mRNA. /FEA=mRNA /GEN=FLJ21276 /PROD=hypothetical protein FLJ21276 /DB\_XREF=gi:13375863 /UG=Hs.41502 hypothetical protein FLJ21276 /FL=gb:NM\_024633.1

209201\_x\_at gb:L01639.1 /DEF=Human (clone HSY3RR) neuropeptide Y receptor (NPYR) mRNA, complete cds. /FEA=mRNA /GEN=NPYR /PROD=neuropeptide Y receptor /DB\_XREF=gi:189313 /UG=Hs.89414 chemokine (C-X-C motif), receptor 4 (fusin) /FL=gb:L01639.1 gb:AF025375.1 gb:M99293.1 gb:L06797.1 gb:NM\_003467.1 gb:AF147204.1

212511\_at Consensus includes gb:AI766247 /FEA=EST /DB\_XREF=gi:5232756 /DB\_XREF=est:wh68f03.x1 /CLONE=IMAGE:2385917 /UG=Hs.7885 phosphatidylinositol binding clathrin assembly protein

207702\_s\_at gb:NM\_012301.1 /DEF=Homo sapiens atrophin-1 interacting protein 1; activin receptor interacting protein 1 (KIAA0705), mRNA. /FEA=mRNA /GEN=KIAA0705 /PROD=atrophin-1 interacting protein 1; activin receptor interacting protein 1 /DB\_XREF=gi:6912461 /UG=Hs.22599 atrophin-1 interacting protein 1; activin receptor interacting protein 1 /FL=gb:AF038563.1 gb:NM\_012301.1

207681\_at gb:NM\_001504.1 /DEF=Homo sapiens G protein-coupled receptor 9 (GPR9), mRNA. /FEA=mRNA /GEN=GPR9 /PROD=G protein-coupled receptor 9 /DB\_XREF=gi:4504098 /UG=Hs.198252 G protein-coupled receptor 9 /FL=gb:NM\_001504.1

216322\_at Consensus includes gb:D28586.1 /DEF=Human mRNA for LFA-3(delta D2), partial cds. /FEA=mRNA /PROD=LFA-3(delta D2) /DB\_XREF=gi:466540 /UG=Hs.75626 CD58 antigen, (lymphocyte function-associated antigen 3)

221973\_at Consensus includes gb:AI983904 /FEA=EST /DB\_XREF=gi:5811123 /DB\_XREF=est:wz55d05.x1 /CLONE=IMAGE:2561961 /UG=Hs.79732 fibulin 1

207092\_at gb:NM\_000230.1 /DEF=Homo sapiens leptin (murine obesity homolog) (LEP), mRNA. /FEA=mRNA /GEN=LEP /PROD=leptin precursor /DB\_XREF=gi:4557714 /UG=Hs.194236 leptin (murine obesity homolog) /FL=gb:U43653.1 gb:AF008123.1 gb:NM\_000230.1 gb:U18915.1 gb:D49487.1

204019\_s\_at gb:NM\_015677.1 /DEF=Homo sapiens hypothetical protein (DKFZP586F1318), mRNA. /FEA=mRNA /GEN=DKFZP586F1318 /PROD=hypothetical protein /DB\_XREF=gi:7661669 /UG=Hs.25213 hypothetical protein /FL=gb:NM\_015677.1

# Affy Probe\_ID

## Sequence Description

212109_at	Consensus includes gb:AI590869 /FEA=EST /DB_XREF=gi:4599917 /DB_XREF=est:tw88g03.x1 /CLONE=IMAGE:2266804 /UG=Hs.172035 hypothetical protein similar to mouse HN1 (Hematological and Neurological expressed sequence 1)
215563_s_at	Consensus includes gb:U28055.1 /DEF=Homo sapiens hepatocyte growth factor-like protein homolog mRNA, partial cds. /FEA=mRNA /PROD=hepatocyte growth factor-like protein homolog /DB_XREF=gi:1141776 /UG=Hs.278657 macrophage stimulating 1 (hepatocyte growth factor-like)
202696_at	gb:Nm_005109.1 /DEF=Homo sapiens oxidative-stress responsive 1 (OSR1), mRNA. /FEA=mRNA /GEN=OSR1 /PROD=oxidative-stress responsive 1 /DB_XREF=gi:4826877 /UG=Hs.95220 oxidative-stress responsive 1 /FL=gb:AB017642.1 gb:Nm_005109.1 gb:AB029024.1
222143_s_at	Consensus includes gb:AY007098.1 /DEF=Homo sapiens clone TCCCIA00183 mRNA sequence. /FEA=mRNA /DB_XREF=gi:9955989 /UG=Hs.27556 hypothetical protein FLJ22405
208161_s_at	gb:Nm_020037.1 /DEF=Homo sapiens ATP-binding cassette, sub-family C (CFTRMRP), member 3 (ABCC3), transcript variant MRP3A, mRNA. /FEA=mRNA /GEN=ABCC3 /PROD=ATP-binding cassette, sub-family C, member 3, isoform MRP3A /DB_XREF=gi:9955971 /UG=Hs.90786 ATP-binding cassette, sub-family C (CFTRMRP), member 3 /FL=gb:AF085691.1 gb:Nm_020037.1
214353_at	Consensus includes gb:AW241864 /FEA=EST /DB_XREF=gi:6575618 /DB_XREF=est:xn76b12.x1 /CLONE=IMAGE:2700383 /UG=Hs.252387 cadherin, EGF LAG seven-pass G-type receptor 1, flamingo (Drosophila) homolog
55616_at	Cluster Incl. AI703342:wd93b03.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2339117 /clone_end=3 /gb=AI703342 /gi=4991242 /ug=Hs.91668 /len=494
208647_at	Consensus includes gb:AA872727 /FEA=EST /DB_XREF=gi:2968849 /DB_XREF=est:ob11c12.s1 /CLONE=IMAGE:1323382 /UG=Hs.48876 farnesyl-diphosphate farnesyltransferase 1 /FL=gb:L06070.1 gb:L06105.1 gb:Nm_004462.1
219659_at	Consensus includes gb:AU146927 /FEA=EST /DB_XREF=gi:11008448 /DB_XREF=est:AU146927 /CLONE=HEMBB1001906 /UG=Hs.130489 ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2 /FL=gb:AF236871.1 gb:Nm_016529.1



# **Affy Probe ID**

## **Sequence Description**

201470\_at gb:NM\_004832.1 /DEF=Homo sapiens glutathione-S-transferase like; glutathione transferase omega (GSTTLP28), mRNA. /FEA=mRNA /GEN=GSTTLP28 /PROD=glutathione-S-transferase like /DB\_XREF=gi:4758483 /UG=Hs.11465 glutathione-S-transferase like; glutathione transferase omega /FL=gb:BC000127.1 gb:U90313.1

209561\_at gb:NM\_004832.1 gb:AF212303.1

214162\_at gb:L38969.1 /DEF=Homo sapiens thrombospondin 3 (THBS3) mRNA, complete cds. /FEA=mRNA /GEN=THBS3 /PROD=thrombospondin 3 /DB\_XREF=gi:886298 /UG=Hs.169875 thrombospondin 3 /FL=gb:NM\_007112.1 gb:L38969.1

204681\_s\_at Consensus includes gb:AF070541.1 /DEF=Homo sapiens clones 24714 and 24715 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:3387900 /UG=Hs.4267 Homo sapiens clones 24714 and 24715 mRNA sequence

203132\_at gb:NM\_012294.1 /DEF=Homo sapiens guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF (KIAA0277), mRNA. /FEA=mRNA /GEN=KIAA0277 /PROD=guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF /DB\_XREF=gi:6912455 /UG=Hs.80620 guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF /FL=gb:D87467.1 gb:NM\_012294.1

202973\_x\_at gb:NM\_000321.1 /DEF=Homo sapiens retinoblastoma 1 (including osteosarcoma) (RB1), mRNA. /FEA=mRNA /GEN=RB1 /PROD=retinoblastoma 1 (including osteosarcoma) /DB\_XREF=gi:4506434 /UG=Hs.75770 retinoblastoma 1 (including osteosarcoma) /FL=gb:M33647.1 gb:M15400.1 gb:M28419.1 gb:NM\_000321.1

214023\_x\_at gb:NM\_014883.1 /DEF=Homo sapiens KIAA0914 gene product (KIAA0914), mRNA. /FEA=mRNA /GEN=KIAA0914 /PROD=KIAA0914 gene product /DB\_XREF=gi:7662375 /UG=Hs.177664 KIAA0914 gene product /FL=gb:AB020721.1 gb:NM\_014883.1

212254\_s\_at Consensus includes gb:AL533838 /FEA=EST /DB\_XREF=gi:12797331 /DB\_XREF=est:AL533838 /CLONE=CS0DF002YB07 (3 prime) /UG=Hs.179661 tubulin, beta polypeptide

218028\_at Consensus includes gb:AI798790 /FEA=EST /DB\_XREF=gi:5364262 /DB\_XREF=est:we92f11.x1 /CLONE=IMAGE:2348589 /UG=Hs.198689 KIAA0728 protein

203263\_s\_at gb:NM\_016031.1 /DEF=Homo sapiens elongation of very long chain fatty acids (FEN1Elo2, yeast)-like 1 (ELOVL1), mRNA. /FEA=mRNA /GEN=ELOVL1 /PROD=elongation of very long chain fatty acids(FEN1Elo2, SUR4Elo3, yeast)-like 1 /DB\_XREF=gi:7705798 /UG=Hs.25597 elongation of very long chain fatty acids (FEN1Elo2, SUR4Elo3, yeast)-like 1 /FL=gb:BC000618.1 gb:NM\_022821.1 gb:AF151846.1 gb:NM\_016031.1

Consensus includes gb:AI625739 /FEA=EST /DB\_XREF=gi:4650670 /DB\_XREF=est:ty65g05.x1 /CLONE=IMAGE:2283992 /UG=Hs.54697 Cdc42 guanine exchange factor (GEF) 9 /FL=gb:NM\_015185.1

**Affy Probe\_ID**

**Sequence Description**

212360_at	Consensus includes gb:AI916249 /FEA=EST /DB_XREF=gi:5636104 /DB_XREF=est:wg99c01.x1 /CLONE=IMAGE:2379360 /UG=Hs.82927 adenosine monophosphate deaminase 2 (isoform L) /FL=gb:NM_004037.2
200732_s_at	Consensus includes gb:AL578310 /FEA=EST /DB_XREF=gi:12942259 /DB_XREF=est:AL578310 /CLONE=CS0DK010YM06 (3 prime) /UG=Hs.227777 protein tyrosine phosphatase type IVA, member 1 /FL=gb:U48296.1 gb:NM_003463.1
203912_s_at	gb:NM_006730.1 /DEF=Homo sapiens deoxyribonuclease I-like 1 (DNASE1L1), mRNA. /FEA=mRNA /GEN=DNASE1L1 /PROD=deoxyribonuclease I-like 1 /DB_XREF=gi:5803006 /UG=Hs.77091 deoxyribonuclease I-like 1 /FL=gb:U06846.1 gb:NM_006730.1
212246_at	Consensus includes gb:BE880828 /FEA=EST /DB_XREF=gi:10329604 /DB_XREF=est:601493261F1 /CLONE=IMAGE:3895407 /UG=Hs.84775 Human transposon-like element mRNA
214204_at	Consensus includes gb:BF224076 /FEA=EST /DB_XREF=gi:11131303 /DB_XREF=est:7q83e08.x1 /CLONE=IMAGE:3704942 /UG=Hs.307438 Human DNA sequence from clone 495O10 on chromosome 6q26-27. Contains an RPL37A (60S Ribosomal Protein L37A) pseudogene, the last exon of a gene for a novel protein similar to worm E04F6.2, ESTs, STSs and GSSs
207643_s_at	gb:NM_001065.1 /DEF=Homo sapiens tumor necrosis factor receptor superfamily, member 1A (TNFRSF1A), mRNA. /FEA=mRNA /GEN=TNFRSF1A /PROD=tumor necrosis factor receptor 1 (55kD) /DB_XREF=gi:4507574 /UG=Hs.159 tumor necrosis factor receptor superfamily, member 1A /FL=gb:NM_001065.1
200611_s_at	gb:AB010427.2 /DEF=Homo sapiens mRNA for NORI-1, complete cds. /FEA=mRNA /PROD=NORI-1 /DB_XREF=gi:5103672 /UG=Hs.85100 WD repeat domain 1 /FL=gb:BC000201.1 gb:BC002489.1 gb:AF020056.1 gb:AB010427.2 gb:NM_017491.1
218153_at	gb:NM_024537.1 /DEF=Homo sapiens hypothetical protein FLJ12118 (FLJ12118), mRNA. /FEA=mRNA /GEN=FLJ12118 /PROD=hypothetical protein FLJ12118 /DB_XREF=gi:13375694 /UG=Hs.7779 hypothetical protein FLJ12118 /FL=gb:NM_024537.1
219627_at	gb:NM_024910.1 /DEF=Homo sapiens hypothetical protein FLJ12700 (FLJ12700), mRNA. /FEA=mRNA /GEN=FLJ12700 /PROD=hypothetical protein FLJ12700 /DB_XREF=gi:13376370 /UG=Hs.250547 hypothetical protein FLJ12700 /FL=gb:NM_024910.1
204333_s_at	gb:NM_000027.1 /DEF=Homo sapiens aspartylglucosaminidase (AGA), mRNA. /FEA=mRNA /GEN=AGA /PROD=aspartylglucosaminidase precursor /DB_XREF=gi:4557272 /UG=Hs.207776 aspartylglucosaminidase /FL=gb:M64073.1 gb:NM_000027.1

# Affy Probe\_ID

## Sequence Description

203782\_s\_at gb:NM\_005035.1 /DEF=Homo sapiens polymerase (RNA) mitochondrial (DNA directed) (POLRMT), mRNA. /FEA=mRNA /GEN=POLRMT /PROD=polymerase (RNA) mitochondrial (DNA directed) /DB\_XREF=gi:4826925 /UG=Hs.153880 polymerase (RNA) mitochondrial (DNA directed) /FL=gb:U75370.1 gb:NM\_005035.1

203435\_s\_at gb:NM\_007287.1 /DEF=Homo sapiens membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) (MME), transcript variant 1bis, mRNA. /FEA=mRNA /GEN=MME /PROD=membrane metallo-endopeptidase /DB\_XREF=gi:6042199 /UG=Hs.1298 membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) /FL=gb:J03779.1 gb:NM\_007287.1 gb:NM\_007288.1

205403\_at gb:NM\_004633.1 /DEF=Homo sapiens interleukin 1 receptor, type II (IL1R2), mRNA. /FEA=mRNA /GEN=IL1R2 /PROD=interleukin 1 receptor, type II /DB\_XREF=gi:4758597 /UG=Hs.25333 interleukin 1 receptor, type II /FL=gb:U74649.1 gb:NM\_004633.1

202719\_s\_at gb:BC001451.1 /DEF=Homo sapiens, testin, clone MGC:1146, mRNA, complete cds. /FEA=mRNA /PROD=testin /DB\_XREF=gi:12655188 /UG=Hs.165986 testin /FL=gb:AF245356.1 gb:AF245357.1 gb:BC001451.1 gb:NM\_015641.1

200701\_at gb:NM\_006432.1 /DEF=Homo sapiens epididymal secretory protein (19.5kD) (HE1), mRNA. /FEA=mRNA /GEN=HE1 /PROD=epididymal secretory protein (19.5kD) /DB\_XREF=gi:5453677 /UG=Hs.119529 epididymal secretory protein (19.5kD) /FL=gb:BC002532.1 gb:NM\_006432.1

208749\_x\_at gb:AF085357.1 /DEF=Homo sapiens flotillin mRNA, complete cds. /FEA=mRNA /PROD=flotillin /DB\_XREF=gi:5114048 /UG=Hs.179986 flotillin 1 /FL=gb:BC001146.1 gb:AF085357.1 gb:AF089750.2 gb:NM\_005803.2

220980\_s\_at gb:NM\_031284.1 /DEF=Homo sapiens hypothetical protein DKFZp434B195 (DKFZP434B195), mRNA. /FEA=mRNA /GEN=DKFZP434B195 /PROD=hypothetical protein DKFZp434B195 /DB\_XREF=gi:13775193 /FL=gb:NM\_031284.1

201126\_s\_at gb:NM\_002406.2 /DEF=Homo sapiens mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT1), mRNA. /FEA=mRNA /GEN=MGAT1 /PROD=mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase /DB\_XREF=gi:6031182 /UG=Hs.151513 mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase /FL=gb:M55621.1 gb:NM\_002406.2

202842\_s\_at gb:AL080081.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F1862 (from clone DKFZp564F1862); complete cds. /FEA=mRNA /GEN=DKFZp564F1862 /PROD=hypothetical protein /DB\_XREF=gi:5262493 /UG=Hs.6790 DnaJ (Hsp40) homolog, subfamily B, member 9 /FL=gb:AF083247.1 gb:AL080081.1 gb:AB026908.1 gb:NM\_012328.1

Affy Probe_ID	Sequence Description
220437_at	gb:NM_018687.1 /DEF=Homo sapiens hepatocellular carcinoma-associated gene TD26 (LOC55908), mRNA. /FEA=mRNA /GEN=LOC55908 /PROD=hepatocellular carcinoma-associated gene TD26 /DB_XREF=gi:8923901 /UG=Hs.224121 hepatocellular carcinoma-associated gene TD26 /FL=gb:NM_018687.1
205363_at	gb:NM_003986.1 /DEF=Homo sapiens butyrobetaine (gamma), 2-oxoglutarate dioxygenase (gamma-butyrobetaine hydroxylase) 1 (BBOX1), mRNA. /FEA=mRNA /GEN=BBOX1 /PROD=gamma-butyrobetaine hydroxylase /DB_XREF=gi:4502368 /UG=Hs.9667 butyrobetaine (gamma), 2-oxoglutarate dioxygenase (gamma-butyrobetaine hydroxylase) 1 /FL=gb:AF082868.1 gb:NM_003986.1
218865_at	gb:NM_022746.1 /DEF=Homo sapiens hypothetical protein FLJ22390 (FLJ22390), mRNA. /FEA=mRNA /GEN=FLJ22390 /PROD=hypothetical protein FLJ22390 /DB_XREF=gi:12232404 /UG=Hs.10974 hypothetical protein FLJ22390 /FL=gb:NM_022746.1
210662_at	gb:BC000879.1 /DEF=Homo sapiens, Similar to kynureninase (L-kynurenine hydrolase), clone MGC:5080, mRNA, complete cds. /FEA=mRNA /PROD=Similar to kynureninase (L-kynureninehydrolase) /DB_XREF=gi:12654128 /UG=Hs.169139 kynureninase (L-kynurenine hydrolase) /FL=gb:BC000879.1
204531_s_at	gb:NM_007295.1 /DEF=Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant BRCA1b, mRNA. /FEA=mRNA /GEN=BRCA1 /PROD=breast cancer 1, early onset /DB_XREF=gi:6552300 /UG=Hs.194143 breast cancer 1, early onset /FL=gb:NM_007295.1
202515_at	Consensus includes gb:BG251175 /FEA=EST /DB_XREF=gi:12760991 /DB_XREF=est:602364982F1 /CLONE=IMAGE:4473165 /UG=Hs.154294 discs, large (Drosophila) homolog 1 /FL=gb:NM_004087.1 gb:U13896.1
37152_at	Cluster Incl. L07592:Human peroxisome proliferator activated receptor mRNA, complete cds /cds=(337,1662) /gb=L07592 /gi=190229 /ug=Hs.106415 /len=3301
216381_x_at	Consensus includes gb:AL035413 /DEF=Human DNA sequence from clone RP4-657E11 on chromosome 1p35.1-36.23 Contains 3 part of the CAPZB (capping protein (actin filament) muscle Z-line, beta) gene, genes for aldo-keto reductase family 7 (aflatoxin aldehyde reductase) members A2 (AKR7A2... /FEA=mRNA_1 /DB_XREF=gi:6010110 /UG=Hs.284236 aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase)
220162_s_at	gb:NM_022352.1 /DEF=Homo sapiens caspase recruitment domain protein 9 (LOC64170), mRNA. /FEA=mRNA /GEN=LOC64170 /PROD=caspase recruitment domain protein 9 /DB_XREF=gi:11641262 /UG=Hs.271815 caspase recruitment domain protein 9 /FL=gb:NM_022352.1

# Affy Probe\_ID

## Sequence Description

219687\_at gb:NM\_018194.1 /DEF=Homo sapiens hypothetical protein FLJ10724 (FLJ10724), mRNA. /FEA=mRNA. /GEN=FLJ10724 /PROD=hypothetical protein FLJ10724 /DB\_XREF=gi:8922620 /UG=Hs.58650 melanoma antigen recognized by T cells 2 /FL=gb:NM\_018194.1

202214\_s\_at gb:NM\_003588.1 /DEF=Homo sapiens cullin 4B (CUL4B), mRNA. /FEA=mRNA /GEN=CUL4B /PROD=cullin 4B /DB\_XREF=gi:13270466 /UG=Hs.155976 cullin 4B /FL=gb:NM\_003588.1 gb:AB014595.1

215346\_at Consensus includes gb:BF664114 /FEA=EST /DB\_XREF=gi:11938009 /DB\_XREF=est:602145911F1 /CLONE=IMAGE:4309610 /UG=Hs.25648 tumor necrosis factor receptor superfamily, member 5

219768\_at gb:NM\_024626.1 /DEF=Homo sapiens hypothetical protein FLJ22418 (FLJ22418), mRNA. /FEA=mRNA /GEN=FLJ22418 /PROD=hypothetical protein FLJ22418 /DB\_XREF=gi:13375849 /UG=Hs.36563 hypothetical protein FLJ22418 /FL=gb:NM\_024626.1

202464\_s\_at gb:NM\_004566.1 /DEF=Homo sapiens 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 (PFKFB3), mRNA. /FEA=mRNA /GEN=PFKFB3 /PROD=6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /DB\_XREF=gi:4758899 /UG=Hs.195471 6-phosphofructo-2-kinasefructose-2,6-biphosphatase 3 /FL=gb:D49817.1 gb:AF109735.1

212762\_s\_at gb:NM\_004566.1

206487\_at Consensus includes gb:AI375916 /FEA=EST /DB\_XREF=gi:4175906 /DB\_XREF=est:tc14c12.x1 /CLONE=IMAGE:2063830 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG-box)

218817\_at gb:NM\_025154.1 /DEF=Homo sapiens KIAA0810 protein (KIAA0810), mRNA. /FEA=mRNA /GEN=KIAA0810 /PROD=hypothetical protein FLJ12407 /DB\_XREF=gi:13435158 /UG=Hs.7531 KIAA0810 protein /FL=gb:NM\_025154.1

205227\_at gb:NM\_021928.1 /DEF=Homo sapiens hypothetical protein FLJ22649 similar to signal peptidase SPC2223 (FLJ22649), mRNA. /FEA=mRNA /GEN=FLJ22649 /PROD=hypothetical protein FLJ22649 similar to signal peptidase SPC2223 /DB\_XREF=gi:11345461 /UG=Hs.42194 hypothetical protein FLJ22649 similar to signal peptidase SPC2223 /FL=gb:NM\_021928.1 gb:AL136660.1

208795\_s\_at gb:NM\_002182.1 /DEF=Homo sapiens interleukin 1 receptor accessory protein (IL1RAP), mRNA. /FEA=mRNA /GEN=IL1RAP /PROD=interleukin 1 receptor accessory protein /DB\_XREF=gi:4504660 /UG=Hs.173880 interleukin 1 receptor accessory protein /FL=gb:AF029213.1 gb:AB006537.1 gb:NM\_002182.1

208795\_s\_at gb:D55716.1 /DEF=Human mRNA for P1cdc47, complete cds. /FEA=mRNA /GEN=P1cdc47 /PROD=P1cdc47 /DB\_XREF=gi:1255616 /UG=Hs.77152 minichromosome maintenance deficient (S. cerevisiae) 7 /FL=gb:D55716.1

# Affy Probe\_ID

## Sequence Description

36129\_at Cluster Incl. AB007857:Homo sapiens KIAA0397 mRNA, complete cds /cds=(54,1520) /gb=AB007857 /gi=2662074 /ug=Hs.7416 /len=6629

201827\_at gb:AF113019.1 /DEF=Homo sapiens PRO2451 mRNA, complete cds. /FEA=mRNA /PROD=PRO2451 /DB\_XREF=gi:6642761 /UG=Hs.250581 SWISNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2 /FL=gb:U66618.1 gb:Nm\_003077.1 gb:AF113019.1

200599\_s\_at gb:Nm\_003299.1 /DEF=Homo sapiens tumor rejection antigen (gp96) 1 (TRA1), mRNA. /FEA=mRNA /GEN=TRA1 /PROD=tumor rejection antigen (gp96) 1 /DB\_XREF=gi:4507676 /UG=Hs.82689 tumor rejection antigen (gp96) 1 /FL=gb:Nm\_003299.1

210473\_s\_at gb:M37712.1 /DEF=Human p58GTA (galactosyltransferase associated protein kinase) mRNA, complete cds. /FEA=mRNA /PROD=p58GTA protein kinase /DB\_XREF=gi:189480 /UG=Hs.183418 cell division cycle 2-like 1 (PITSLRE proteins) /FL=gb:M37712.1 gb:U04819.1

207034\_s\_at gb:Nm\_030379.1 /DEF=Homo sapiens GLI-Kruppel family member GLI2 (GLI2), transcript variant 1, mRNA. /FEA=mRNA /GEN=GLI2 /PROD=GLI-Kruppel family member GLI2, isoform alpha /DB\_XREF=gi:13518230 /UG=Hs.111867 GLI-Kruppel family member GLI2 /FL=gb:Nm\_030379.1 gb:AB007295.1

212875\_s\_at Consensus includes gb:AP001745 /DEF=Homo sapiens genomic DNA, chromosome 21q, section 89105 /FEA=mRNA\_3 /DB\_XREF=gi:7768737 /UG=Hs.16007 chromosome 21 open reading frame 25

201793\_x\_at Consensus includes gb:BF509099 /FEA=EST /DB\_XREF=gi:11592397 /DB\_XREF=est:UI-H-BI4-aou-e-05-0-UI.sl /CLONE=IMAGE:3086288 /UG=Hs.15087 KIAA0250 gene product /FL=gb:D87437.1 gb:Nm\_014837.1

217992\_s\_at gb:Nm\_024329.1 /DEF=Homo sapiens hypothetical protein MGC4342 (MGC4342), mRNA. /FEA=mRNA /GEN=MGC4342 /PROD=hypothetical protein MGC4342 /DB\_XREF=gi:13443015 /UG=Hs.301342 hypothetical protein MGC4342 /FL=gb:BC003033.1 gb:Nm\_024329.1

211731\_x\_at gb:BC005904.1 /DEF=Homo sapiens, synovial sarcoma, X breakpoint 3, clone MGC:14495, mRNA, complete cds. /FEA=mRNA /PROD=synovial sarcoma, X breakpoint 3 /DB\_XREF=gi:13543493 /FL=gb:BC005904.1

219408\_at gb:Nm\_019023.1 /DEF=Homo sapiens hypothetical protein (FLJ10640), mRNA. /FEA=mRNA /GEN=FLJ10640 /PROD=hypothetical protein /DB\_XREF=gi:9506614 /UG=Hs.91753 hypothetical protein /FL=gb:BC000146.1 gb:Nm\_019023.1

218506\_x\_at gb:Nm\_018459.1 /DEF=Homo sapiens uncharacterized bone marrow protein BM045 (BM045), mRNA. /FEA=mRNA /GEN=BM045 /PROD=uncharacterized bone marrow protein BM045 /DB\_XREF=gi:8922103 /UG=Hs.8750 uncharacterized bone marrow protein BM045 /FL=gb:AF217521.1 gb:Nm\_018459.1

# Affy Probe ID

## Sequence Description

200742\_s\_at Consensus includes gb:BG231932 /FEA=EST /DB\_XREF=gi:12727071 /DB\_XREF=est:naf34b12.x1 /CLONE=IMAGE:4142926 /UG=Hs.20478 ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease) /FL=gb:AF017456.1 gb:NM\_000391.2

218584\_at gb:NM\_024549.1 /DEF=Homo sapiens hypothetical protein FLJ21127 (FLJ21127), mRNA. /FEA=mRNA /GEN=FLJ21127 /PROD=hypothetical protein FLJ21127 /DB\_XREF=gi:13375712 /UG=Hs.11500 hypothetical protein FLJ21127 /FL=gb:NM\_024549.1

205026\_at gb:NM\_012448.1 /DEF=Homo sapiens signal transducer and activator of transcription 5B (STAT5B), mRNA. /FEA=mRNA /GEN=STAT5B /PROD=signal transducer and activator of transcription5B /DB\_XREF=gi:6912687 /UG=Hs.244613 signal transducer and activator of transcription 5B /FL=gb:U47686.1 gb:NM\_012448.1 gb:U48730.2

204861\_s\_at gb:NM\_004536.1 /DEF=Homo sapiens baculoviral IAP repeat-containing 1 (BIRC1), mRNA. /FEA=mRNA /GEN=BIRC1 /PROD=baculoviral IAP repeat-containing 1 /DB\_XREF=gi:4758751 /UG=Hs.79019 baculoviral IAP repeat-containing 1 /FL=gb:U19251.1 gb:NM\_004536.1

210512\_s\_at gb:AF022375.1 /DEF=Homo sapiens vascular endothelial growth factor mRNA, complete cds. /FEA=mRNA /PROD=vascular endothelial growth factor /DB\_XREF=gi:3719220 /UG=Hs.73793 vascular endothelial growth factor /FL=gb:M32977.1 gb:AF022375.1 gb:NM\_003376.1 gb:AB021221.1 gb:AF091352.1

209019\_s\_at gb:AF316873.1 /DEF=Homo sapiens protein kinase BRPK mRNA, complete cds. /FEA=mRNA /PROD=protein kinase BRPK /DB\_XREF=gi:13492051 /UG=Hs.6163 Homo sapiens protein kinase BRPK mRNA, complete cds /FL=gb:AF316873.1

220936\_s\_at gb:NM\_018267.1 /DEF=Homo sapiens hypothetical protein FLJ10903 (FLJ10903), mRNA. /FEA=mRNA /GEN=FLJ10903 /PROD=hypothetical protein FLJ10903 /DB\_XREF=gi:8922757 /UG=Hs.36727 hypothetical protein FLJ10903 /FL=gb:NM\_018267.1

218729\_at gb:NM\_020169.1 /DEF=Homo sapiens latexin protein (LXN), mRNA. /FEA=mRNA /GEN=LXN /PROD=latexin protein /DB\_XREF=gi:9910395 /UG=Hs.109276 latexin protein /FL=gb:BC005346.1 gb:AF282626.1 gb:NM\_020169.1

213940\_s\_at Consensus includes gb:AU145053 /FEA=EST /DB\_XREF=gi:11006574 /DB\_XREF=est:AU145053 /CLONE=HEMBA1003755 /UG=Hs.301763 KIAA0554 protein

220453\_at gb:NM\_017765.1 /DEF=Homo sapiens hypothetical protein FLJ20320 (FLJ20320), mRNA. /FEA=mRNA /GEN=FLJ20320 /PROD=hypothetical protein FLJ20320 /DB\_XREF=gi:8923300 /UG=Hs.263081 hypothetical protein FLJ20320 /FL=gb:NM\_017765.1

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# Affy Probe ID

## Sequence Description

212976_at	Consensus includes gb:R41498 /FEA=EST /DB_XREF=gi:816800 /DB_XREF=est:yf86h08.s1 /CLONE=IMAGE:29486 /UG=Hs.199243 KIAA0231 protein
222159_at	Consensus includes gb:AK023058.1 /DEF=Homo sapiens cDNA FLJ12996 fis, clone NT2RP3000235. /FEA=mRNA /DB_XREF=gi:10434801 /UG=Hs.287546 Homo sapiens cDNA FLJ12996 fis, clone NT2RP3000235
212055_at	Consensus includes gb:BF689173 /FEA=EST /DB_XREF=gi:11974581 /DB_XREF=est:602184834T1 /CLONE=IMAGE:4299201 /UG=Hs.22981 DKFZP586M1523 protein
219024_at	gb:NM_021622.1 /DEF=Homo sapiens pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1 (PLEKHA1), mRNA. /FEA=mRNA /GEN=PLEKHA1 /PROD=pleckstrin homology domain-containing, family A(phosphoinositide binding specific) member 1 /DB_XREF=gi:11055985 /UG=Hs.17757 pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1 /FL=gb:AF286160.1
205595_at	gb:NM_021622.1 gb:BC001136.1 gb:NM_001944.1 /DEF=Homo sapiens desmoglein 3 (pemphigus vulgaris antigen) (DSG3), mRNA. /FEA=mRNA /GEN=DSG3 /PROD=desmoglein 3 preproprotein /DB_XREF=gi:4503404 /UG=Hs.1925 desmoglein 3 (pemphigus vulgaris antigen) /FL=gb:M76482.1 gb:NM_001944.1
203509_at	gb:NM_003105.2 /DEF=Homo sapiens sortilin-related receptor, L(DLR class) A repeats-containing (SORL1), mRNA. /FEA=mRNA /GEN=SORL1 /PROD=sortilin-related receptor, L(DLR class) Arepeats-containing /DB_XREF=gi:6325473 /UG=Hs.278571 sortilin-related receptor, L(DLR class) A repeats-containing /FL=gb:U60975.2 gb:NM_003105.2
208877_at	Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=est:zd75f10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF092132.1
212350_at	Consensus includes gb:AB029031.1 /DEF=Homo sapiens mRNA for KIAA1108 protein, partial cds. /FEA=mRNA /GEN=KIAA1108 /PROD=KIAA1108 protein /DB_XREF=gi:5689552 /UG=Hs.278586 KIAA1108 protein
222076_at	Consensus includes gb:BE222436 /FEA=EST /DB_XREF=gi:8909754 /DB_XREF=est:hv90h11.x1 /CLONE=IMAGE:3180741 /UG=Hs.799 diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)
206628_at	gb:NM_000343.1 /DEF=Homo sapiens solute carrier family 5 (sodiumglucose cotransporter), member 1 (SLC5A1), mRNA. /FEA=mRNA /GEN=SLC5A1 /PROD=solute carrier family 5 (sodiumglucosecotransporter), member 1 /DB_XREF=gi:4507030 /UG=Hs.1964 solute carrier family 5 (sodiumglucose cotransporter), member 1 /FL=gb:M24847.1 gb:NM_000343.1



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220148_at	gb:NM_022568.1 /DEF=Homo sapiens aldehyde dehydrogenase 12 (ALDH12), mRNA. /FEA=mRNA /GEN=ALDH12 /PROD=aldehyde dehydrogenase 12 /DB_XREF=gi:12007647 /UG=Hs.18443 aldehyde dehydrogenase 8 family, member A1 /FL=gb:AF303134.1 gb:NM_022568.1
218687_s_at	gb:NM_017648.1 /DEF=Homo sapiens hypothetical protein FLJ20063 (FLJ20063), mRNA. /FEA=mRNA /GEN=FLJ20063 /PROD=hypothetical protein FLJ20063 /DB_XREF=gi:8923068 /UG=Hs.5940 hypothetical protein FLJ20063 /FL=gb:AB035807.1 gb:NM_017648.1
202543_s_at	gb:BC005359.1 /DEF=Homo sapiens, glia maturation factor, beta, clone MGC:12462, mRNA, complete cds. /FEA=mRNA /PROD=glia maturation factor, beta /DB_XREF=gi:13529184 /UG=Hs.151413 glia maturation factor, beta /FL=gb:BC005359.1 gb:M86492.1 gb:AB001106.1 gb:NM_004124.1
204491_at	Consensus includes gb:R40917 /FEA=EST /DB_XREF=gi:823119 /DB_XREF=est:yf75d09.s1 /CLONE=IMAGE:28220 /UG=Hs.172081 phosphodiesterase 4D, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E3) /FL=gb:L20969.1 gb:U02882.1 gb:NM_006203.1
212503_s_at	Consensus includes gb:N22859 /FEA=EST /DB_XREF=gi:1137009 /DB_XREF=est:yx56d09.s1 /CLONE=IMAGE:265745 /UG=Hs.227716 KIAA0934 protein
216921_s_at	Consensus includes gb:X90763.1 /DEF=Homo sapiens mRNA for type I keratin. /FEA=mRNA /GEN=hHa5 /PROD=hHa5 hair keratin type I intermediate filament /DB_XREF=gi:1668743 /UG=Hs.73082 keratin, hair, acidic, 5
217664_at	Consensus includes gb:AA780524 /FEA=EST /DB_XREF=gi:2839855 /DB_XREF=est:ac71f01.s1 /CLONE=IMAGE:868057 /UG=Hs.294072 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAMILY J
211997_x_at	SEQUENCE CONTAMINATION WARNING ENTRY H.sapiens
211997_x_at	Consensus includes gb:NM_005324.1 /DEF=Homo sapiens H3 histone, family 3B (H3.3B) (H3F3B), mRNA. /FEA=CDS /GEN=H3F3B /PROD=H3 histone, family 3B (H3.3B) /DB_XREF=gi:4885384 /UG=Hs.180877 H3 histone, family 3B (H3.3B) /FL=gb:NM_005324.1
220233_at	gb:NM_024907.1 /DEF=Homo sapiens hypothetical protein FLJ11798 (FLJ11798), mRNA. /FEA=mRNA /GEN=FLJ11798 /PROD=hypothetical protein FLJ11798 /DB_XREF=gi:13376364 /UG=Hs.249727 hypothetical protein FLJ11798 /FL=gb:NM_024907.1
221297_at	gb:NM_018654.1 /DEF=Homo sapiens G protein-coupled receptor, family C, group 5, member D (GPC5D), mRNA. /FEA=CDS /GEN=GPC5D /PROD=G protein-coupled receptor, family C, group 5, member D /DB_XREF=gi:8923704 /UG=Hs.283073 G protein-coupled receptor, family C, group 5, member D /FL=gb:AF209923.1 gb:NM_018654.1

# Affy Probe\_ID

## Sequence Description

202864\_s\_at gb:NM\_003113.1 /DEF=Homo sapiens nuclear antigen Sp100 (SP100), mRNA. /FEA=mRNA /GEN=SP100 /PROD=nuclear antigen Sp100 /DB\_XREF=gi:4507164 /UG=Hs.77617 nuclear antigen Sp100 /FL=gb:M60618.1 gb:NM\_003113.1

218898\_at gb:NM\_024792.1 /DEF=Homo sapiens hypothetical protein FLJ22282 (FLJ22282), mRNA. /FEA=mRNA /GEN=FLJ22282 /PROD=hypothetical protein FLJ22282 /DB\_XREF=gi:13376161 /UG=Hs.154396 hypothetical protein FLJ22282 /FL=gb:NM\_024792.1

209377\_s\_at gb:AF274949.1 /DEF=Homo sapiens PNAS-24 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-24 /DB\_XREF=gi:12751062 /UG=Hs.77558 thyroid hormone receptor interactor 7 /FL=gb:AF274949.1

209485\_s\_at Consensus includes gb:W19983 /FEA=EST /DB\_XREF=gi:1295853 /DB\_XREF=est:zb38d10.r1 /CLONE=IMAGE:305875 /UG=Hs.252716 Homo sapiens oxysterol-binding protein-related protein (ORP1) mRNA, complete cds /FL=gb:AF274714.1

211612\_s\_at gb:U62858.1 /DEF=Human interleukin-13 receptor mRNA, complete cds. /FEA=mRNA /PROD=interleukin-13 receptor /DB\_XREF=gi:1695875 /FL=gb:U62858.1

210608\_s\_at gb:BC001899.1 /DEF=Homo sapiens, Similar to fucosyltransferase 2 (secretor status included), clone MGC:2062, mRNA, complete cds. /FEA=mRNA /PROD=Similar to fucosyltransferase 2 (secretor status included) /DB\_XREF=gi:12804894 /UG=Hs.46328 fucosyltransferase 2 (secretor status included) /FL=gb:BC001899.1

212335\_at Consensus includes gb:AW167793 /FEA=EST /DB\_XREF=gi:6399401 /DB\_XREF=est:xg56d07.x1 /CLONE=IMAGE:2632333 /UG=Hs.164036 Homo sapiens AKAP350C mRNA sequence, alternatively spliced

212964\_at Consensus includes gb:AI912206 /FEA=EST /DB\_XREF=gi:5632061 /DB\_XREF=est:wd71e08.x1 /CLONE=IMAGE:2337062 /UG=Hs.109445 KIAA1020 protein

221486\_at gb:AF067170.1 /DEF=Homo sapiens alpha endosulfine mRNA, complete cds. /FEA=mRNA /PROD=alpha endosulfine /DB\_XREF=gi:4894373 /UG=Hs.111680 endosulfine alpha /FL=gb:AF067170.1 gb:AF157510.1

208760\_at Consensus includes gb:AL031714 /DEF=Human DNA sequence from clone LA16-358B7 on chromosome 16 Contains the UBE21 gene for ubiquitin-conjugating enzyme E2I (homologous to yeast UBC9), and an RPS20 (40S Ribosomal protein S20) pseudogene. Contains ESTs, STSs. GSSs and a putative CpG is... /FEA=mRNA /DB\_XREF=gi:4775608 /UG=Hs.84285 ubiquitin-conjugating enzyme E2I (homologous to yeast UBC9) /FL=gb:U45328.1 gb:U31933.1 gb:BC000427.1 gb:BC004429.1 gb:U31882.1 gb:U66818.1 gb:U66867.1 gb:U38785.1 gb:NM\_003345.1 gb:U29092.1

# Affy Probe\_ID

## Sequence Description

218284\_at gb:NM\_015400.1 /DEF=Homo sapiens DKFZP586N0721 protein (DKFZP586N0721), mRNA. /FEA=mRNA /GEN=DKFZP586N0721 /PROD=DKFZP586N0721 protein /DB\_XREF=gi:7661691 /UG=Hs.99843 DKFZP586N0721 protein /FL=gb:BC000414.1 gb:AF151027.1 gb:NM\_015400.1

49452\_at Cluster Incl. AI057637:oy31h06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1667483 /clone\_end=3 /gb=AI057637 /gi=3331503 /ug=Hs.21305 /len=599

212579\_at Consensus includes gb:AB014550.1 /DEF=Homo sapiens mRNA for KIAA0650 protein, partial cds. /FEA=mRNA /GEN=KIAA0650 /PROD=KIAA0650 protein /DB\_XREF=gi:3327113 /UG=Hs.8118 KIAA0650 protein

217668\_at Consensus includes gb:BF511164 /FEA=EST /DB\_XREF=gi:11594462 /DB\_XREF=est:UI-H-BI4-aoi-a-09-0-UI.s1 /CLONE=IMAGE:3084952 /UG=Hs.293309 ESTs

217733\_s\_at gb:NM\_021103.1 /DEF=Homo sapiens thymosin, beta 10 (TMSB10), mRNA. /FEA=mRNA /GEN=TMSB10 /PROD=thymosin, beta 10 /DB\_XREF=gi:10863894 /UG=Hs.76293 thymosin, beta 10 /FL=gb:NM\_021103.1 gb:M92381.1 gb:M20259.1

211995\_x\_at Consensus includes gb:AL567820 /FEA=EST /DB\_XREF=gi:12921560 /DB\_XREF=est:AL567820 /CLONE=CS0DF033YD24 (3 prime) /UG=Hs.14376 actin, gamma 1

209853\_s\_at gb:BC002684.1 /DEF=Homo sapiens, proteaseome (prosomal, macropain) 28 subunit, 3, clone MGC:3572, mRNA, complete cds. /FEA=mRNA /PROD=proteaseome (prosomal, macropain) 28 subunit, 3 /DB\_XREF=gi:12803696 /UG=Hs.152978 proteaseome (prosomal, macropain) activator subunit 3 (PA28 gamma; Ki) /FL=gb:BC001423.1 gb:BC002684.1

219079\_at gb:NM\_016230.1 /DEF=Homo sapiens flavohemoprotein b5+b5R (LOC511167), mRNA. /FEA=mRNA /GEN=LOC511167 /PROD=flavohemoprotein b5+b5R /DB\_XREF=gi:7705898 /UG=Hs.5741 flavohemoprotein b5+b5R /FL=gb:AF169803.1 gb:NM\_016230.1

218657\_at gb:NM\_016339.1 /DEF=Homo sapiens Link guanine nucleotide exchange factor II (LOC511195), mRNA. /FEA=mRNA /GEN=LOC511195 /PROD=Link guanine nucleotide exchange factor II /DB\_XREF=gi:7705938 /UG=Hs.118562 Link guanine nucleotide exchange factor II /FL=gb:AF117946.1 gb:NM\_016339.1

205430\_at Consensus includes gb:AL133386 /DEF=Human DNA sequence from clone RP1-181C24 on chromosome 6p11.1-12.2. Contains the 3 end of the BMP5 gene for bone morphogenetic protein 5, ESTs, STSs and GSSs /FEA=mRNA /DB\_XREF=gi:6807532 /UG=Hs.1104 bone morphogenetic protein 5 /FL=gb:NM\_021073.1 gb:M60314.1

# Affy Probe ID

## Sequence Description

201065\_s\_at gb:NM\_001518.1 /DEF=Homo sapiens general transcription factor II, i (GTF2I), mRNA. /FEA=mRNA /GEN=GTF2I /PROD=general transcription factor II, i /DB\_XREF=gi:4504202 /UG=Hs.278589 general transcription factor II, i /FL=gb:U77948.1 gb:AF015553.1 gb:AF038969.1 gb:NM\_001518.1

203638\_s\_at gb:NM\_022969.1 /DEF=Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 2, mRNA. /FEA=mRNA /GEN=FGFR2 /PROD=fibroblast growth factor receptor 2, isoform 2precursor /DB\_XREF=gi:13186252 /UG=Hs.278581 fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) /FL=gb:NM\_022969.1 gb:M97193.1 gb:M80634.1

200884\_at gb:NM\_001823.1 /DEF=Homo sapiens creatine kinase, brain (CKB), mRNA. /FEA=mRNA /GEN=CKB /PROD=creatine kinase, brain /DB\_XREF=gi:4502850 /UG=Hs.173724 creatine kinase, brain /FL=gb:L47647.1 gb:BC001190.1 gb:BC004914.1 gb:M16364.1 gb:M16451.1 gb:NM\_001823.1

221882\_s\_at Consensus includes gb:AI636233 /FEA=EST /DB\_XREF=gi:4687563 /DB\_XREF=est:tz93b07.x1 /CLONE=IMAGE:2296117 /UG=Hs.288940 five-span transmembrane protein M83

211737\_x\_at gb:BC005916.1 /DEF=Homo sapiens, pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1), clone MGC:14512, mRNA, complete cds. /FEA=mRNA /PROD=pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1) /DB\_XREF=gi:13543514 /FL=gb:BC005916.1

203112\_s\_at gb:NM\_005663.1 /DEF=Homo sapiens Wolf-Hirschhorn syndrome candidate 2 (WHSC2), mRNA. /FEA=mRNA /GEN=WHSC2 /PROD=WHSC2 protein /DB\_XREF=gi:5032226 /UG=Hs.21771 Wolf-Hirschhorn syndrome candidate 2 /FL=gb:BC002764.1 gb:AF101434.1 gb:NM\_005663.1

212387\_at Consensus includes gb:BG495771 /FEA=EST /DB\_XREF=gi:13457287 /DB\_XREF=est:602540396F1 /CLONE=IMAGE:4671599 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1000272

209574\_s\_at Consensus includes gb:AI349506 /FEA=EST /DB\_XREF=gi:4086712 /DB\_XREF=est:qp71g08.x1 /CLONE=IMAGE:1928510 /UG=Hs.153498 chromosome 18 open reading frame 1 /FL=gb:AF009424.1

218471\_s\_at gb:NM\_024649.1 /DEF=Homo sapiens hypothetical protein FLJ23590 (FLJ23590), mRNA. /FEA=mRNA /GEN=FLJ23590 /PROD=hypothetical protein FLJ23590 /DB\_XREF=gi:13375892 /UG=Hs.54890 hypothetical protein FLJ23590 /FL=gb:NM\_024649.1

215380\_s\_at Consensus includes gb:AK021779.1 /DEF=Homo sapiens cDNA FLJ11717 fis, clone HEMBA1005241. /FEA=mRNA /DB\_XREF=gi:10433033 /UG=Hs.288649 hypothetical protein MGC3077

Affy Probe_ID	Sequence Description
207135_at	gb:NM_000621.1 /DEF=Homo sapiens 5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA. /FEA=mRNA /GEN=HTR2A /PROD=5-hydroxytryptamine (serotonin) receptor 2A /DB_XREF=gi:10835174 /UG=Hs.298623 5-hydroxytryptamine (serotonin) receptor 2A /FL=gb:NM_000621.1
201537_s_at	gb:BC002682.1 /DEF=Homo sapiens, dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related), clone MGC:3615, mRNA, complete cds. /FEA=mRNA /PROD=dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) /DB_XREF=gi:12803692 /UG=Hs.181046 dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) /FL=gb:BC002682.1 gb:L05147.1 gb:NM_004090.1
219665_at	gb:NM_024815.1 /DEF=Homo sapiens hypothetical protein FLJ22494 (FLJ22494), mRNA. /FEA=mRNA /GEN=FLJ22494 /PROD=hypothetical protein FLJ22494 /DB_XREF=gi:13376205 /UG=Hs.170132 hypothetical protein FLJ22494 /FL=gb:NM_024815.1
206746_at	gb:NM_001195.2 /DEF=Homo sapiens beaded filament structural protein 1, filensin (BFSP1), mRNA. /FEA=mRNA /GEN=BFSP1 /PROD=filensin /DB_XREF=gi:13518027 /UG=Hs.129702 beaded filament structural protein 1, filensin /FL=gb:Y16717.2 gb:NM_001195.2 gb:AF039655.1
214239_x_at	Consensus includes gb:AI560455 /FEA=EST /DB_XREF=gi:4510796 /DB_XREF=est:tn11b01.x1 /CLONE=IMAGE:2167273 /UG=Hs.184669 zinc finger protein 144 (Mel-18)
201096_s_at	Consensus includes gb:AL537042 /FEA=EST /DB_XREF=gi:12800535 /DB_XREF=est:AL537042 /CLONE=CS0DF017YF17 (5 prime) /UG=Hs.75290 ADP-ribosylation factor 4 /FL=gb:BC003364.1 gb:M36341.1 gb:NM_001660.2
31845_at	Cluster Incl. U32645:Human myeloid elf-1 like factor (MEF) mRNA, complete cds /cds=(382,2373) /gb=U32645 /gi=1761934 /ug=Hs.151139 /len=4167
205453_at	gb:NM_002145.1 /DEF=Homo sapiens homeo box B2 (HOXB2), mRNA. /FEA=mRNA /GEN=HOXB2 /PROD=homeo box B2 /DB_XREF=gi:4504464 /UG=Hs.2733 homeo box B2 /FL=gb:NM_002145.1
205265_s_at	gb:NM_005876.1 /DEF=Homo sapiens nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury (APEG1), mRNA. /FEA=mRNA /GEN=APEG1 /PROD=nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury /DB_XREF=gi:5031588 /UG=Hs.21639 nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury /FL=gb:U57099.1 gb:NM_005876.1

Affy Probe_ID	Sequence Description
219369_s_at	gb:NM_023112.1 /DEF=Homo sapiens hypothetical protein FLJ21916 (FLJ21916), mRNA. /FEA=mRNA /GEN=FLJ21916 /PROD=hypothetical protein FLJ21916 /DB_XREF=gi:12962938 /UG=Hs.90034 hypothetical protein FLJ21916 /FL=gb:BC000208.1 gb:NM_023112.1
212569_at	Consensus includes gb:AV699744 /FEA=EST /DB_XREF=gi:10301715 /DB_XREF=est:AV699744 /CLONE=GKCEDF05 /UG=Hs.8118 KIAA0650 protein
219127_at	gb:NM_024320.1 /DEF=Homo sapiens hypothetical protein MGC11242 (MGC11242), mRNA. /FEA=mRNA /GEN=MGC11242 /PROD=hypothetical protein MGC11242 /DB_XREF=gi:13236560 /UG=Hs.36529 hypothetical protein MGC11242 /FL=gb:BC002865.1 gb:NM_024320.1
213169_at	Consensus includes gb:BG109855 /FEA=EST /DB_XREF=gi:12603361 /DB_XREF=est:602279521F1 /CLONE=IMAGE:4367308 /UG=Hs.49476 Homo sapiens clone TUA8 Cri-du-chat region mRNA
216617_s_at	Consensus includes gb:X98405.1 /DEF=H.sapiens mRNA for myelin associated glycoprotein, S-MAG. /FEA=mRNA /GEN=S-MAG /PROD=myelin associated glycoprotein /DB_XREF=gi:1403054 /UG=Hs.1780 myelin associated glycoprotein
202082_s_at	Consensus includes gb:AV748469 /FEA=EST /DB_XREF=gi:10906317 /DB_XREF=est:AV748469 /CLONE=NPCAXA07 /UG=Hs.75232 SEC14 (S. cerevisiae)-like 1 /FL=gb:D67029.1 gb:NM_003003.1
204791_at	gb:NM_003297.1 /DEF=Homo sapiens nuclear receptor subfamily 2, group C, member 1 (NR2C1), mRNA. /FEA=mRNA /GEN=NR2C1 /PROD=nuclear receptor subfamily 2, group C, member 1 /DB_XREF=gi:4507672 /UG=Hs.108301 nuclear receptor subfamily 2, group C, member 1 /FL=gb:M29960.1 gb:NM_003297.1
53202_at	Cluster Incl. AA402435:zt60g10.r1 Homo sapiens cDNA, 5' end /clone=IMAGE-726786 /clone_end=5 /gb=AA402435 /gi=2057299 /ug=Hs.59594 /len=529
203045_at	gb:NM_004148.1 /DEF=Homo sapiens ninjurin 1 (NINJ1), mRNA. /FEA=mRNA /GEN=NINJ1 /PROD=ninjurin 1 /DB_XREF=gi:4758809 /UG=Hs.11342 ninjurin 1 /FL=gb:BC004440.1 gb:U72661.1 gb:U91512.1 gb:NM_004148.1
208891_at	gb:BC003143.1 /DEF=Homo sapiens, dual specificity phosphatase 6, clone MGC:3789, mRNA, complete cds. /FEA=mRNA /PROD=dual specificity phosphatase 6 /DB_XREF=gi:13111942 /UG=Hs.180383 dual specificity phosphatase 6 /FL=gb:BC003562.1 gb:BC003143.1 gb:BC005047.1 gb:AB013382.1 gb:NM_001946.1
219497_s_at	gb:NM_022893.1 /DEF=Homo sapiens B-cell CLL/lymphoma 11A (zinc finger protein) (BCL11A), mRNA. /FEA=mRNA /GEN=BCL11A /PROD=B-cell CLL/lymphoma 11A (zinc finger protein) /DB_XREF=gi:13375629 /UG=Hs.130881 B-cell CLL/lymphoma 11A (zinc finger protein) /FL=gb:NM_022893.1 gb:NM_018014.1

# Affy Probe ID

## Sequence Description

220221\_at gb:NM\_018156.1 /DEF=Homo sapiens hypothetical protein FLJ10619 (FLJ10619), mRNA. /FEA=mRNA /GEN=FLJ10619 /PROD=hypothetical protein FLJ10619 /DB\_XREF=gi:8922552 /UG=Hs.191436 hypothetical protein FLJ10619 /FL=gb:NM\_018156.1  
213636\_at Consensus includes gb:AB028968.1 /DEF=Homo sapiens mRNA for KIAA1045 protein, partial cds. /FEA=mRNA /GEN=KIAA1045 /PROD=KIAA1045 protein /DB\_XREF=gi:5689426 /UG=Hs.7989 KIAA1045 protein  
219067\_s\_at gb:NM\_017615.1 /DEF=Homo sapiens hypothetical protein FLJ20003 (FLJ20003), mRNA. /FEA=mRNA /GEN=FLJ20003 /PROD=hypothetical protein FLJ20003 /DB\_XREF=gi:89233008 /UG=Hs.258798 hypothetical protein FLJ20003 /FL=gb:NM\_017615.1  
216776\_at Consensus includes gb:AK025206.1 /DEF=Homo sapiens cDNA: FLJ21553 fis, clone COL06329. /FEA=mRNA /DB\_XREF=gi:10437671 /UG=Hs.306786 Homo sapiens cDNA: FLJ21553 fis, clone COL06329  
212978\_at Consensus includes gb:AU146004 /FEA=EST /DB\_XREF=gi:11007525 /DB\_XREF=est:AU146004 /CLONE=HEMBA1006523 /UG=Hs.199243 KIAA0231 protein  
201681\_s\_at Consensus includes gb:AB011155.1 /DEF=Homo sapiens mRNA for KIAA0583 protein, partial cds. /FEA=mRNA /GEN=KIAA0583 /PROD=KIAA0583 protein /DB\_XREF=gi:3043689 /UG=Hs.170290 discs, large (Drosophila) homolog 5 /FL=gb:U61843.1 gb:NM\_004747.1  
38037\_at Cluster Incl. M60278:Human heparin-binding EGF-like growth factor mRNA, complete cds /cds=(261,887) /gb=M60278 /gi=183866 /ug=Hs.799 /len=2342  
212693\_at Consensus includes gb:BE670928 /FEA=EST /DB\_XREF=gi:10031469 /DB\_XREF=est:7e43b11.x1 /CLONE=IMAGE:3285213 /UG=Hs.76730 KIAA0301 protein  
220333\_at gb:NM\_017705.1 /DEF=Homo sapiens hypothetical protein FLJ20190 (FLJ20190), mRNA. /FEA=mRNA /GEN=FLJ20190 /PROD=hypothetical protein FLJ20190 /DB\_XREF=gi:8923182 /UG=Hs.257511 hypothetical protein FLJ20190 /FL=gb:NM\_017705.1  
221438\_s\_at gb:NM\_031275.1 /DEF=Homo sapiens testis expressed sequence 12 (TEX12), mRNA. /FEA=CDS /GEN=TEX12 /PROD=testis expressed sequence 12 /DB\_XREF=gi:13775181 /FL=gb:NM\_031275.1  
200989\_at gb:NM\_001530.1 /DEF=Homo sapiens hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA. /FEA=mRNA /GEN=HIF1A /PROD=hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) /DB\_XREF=gi:4504384 /UG=Hs.197540 hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) /FL=gb:U29165.1 gb:AF304431.1 gb:NM\_001530.1 gb:AF207601.1 gb:AF207602.1 gb:U22431.1

**Affy Probe ID**

**Sequence Description**

201397\_at gb:NM\_006623.1 /DEF=Homo sapiens phosphoglycerate dehydrogenase (PHGDH), mRNA. /FEA=mRNA /GEN=PHGDH /PROD=phosphoglycerate dehydrogenase /DB\_XREF=gi:5729973 /UG=Hs.3343 phosphoglycerate dehydrogenase /FL=gb:BC000303.1 gb:BC001349.1 gb:AF006043.1 gb:Nm\_006623.1 gb:AF171237.1

201583\_s\_at gb:NM\_006363.1 /DEF=Homo sapiens Sec23 (S. cerevisiae) homolog B (SEC23B), mRNA. /FEA=mRNA /GEN=SEC23B /PROD=Sec23 (S. cerevisiae) homolog B /DB\_XREF=gi:5454043 /UG=Hs.173497 Sec23 (S. cerevisiae) homolog B /FL=gb:BC005404.1 gb:Nm\_006363.1

203297\_s\_at Consensus includes gb:BG029530 /FEA=EST /DB\_XREF=gi:12418626 /DB\_XREF=est:602297090F1 /CLONE=IMAGE:4391375 /UG=Hs.40154 jumonji (mouse) homolog /FL=gb:Nm\_004973.2 gb:U57592.1

208959\_s\_at gb:BC005374.1 /DEF=Homo sapiens, Similar to RIKEN cDNA 1110001E24 gene, clone MGC:12490, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RIKEN cDNA 1110001E24 gene /DB\_XREF=gi:13529223 /UG=Hs.154023 KIAA0573 protein /FL=gb:BC005374.1

204168\_at gb:NM\_002413.1 /DEF=Homo sapiens microsomal glutathione S-transferase 2 (MGST2), mRNA. /FEA=mRNA /GEN=MGST2 /PROD=microsomal glutathione S-transferase 2 /DB\_XREF=gi:4505180 /UG=Hs.81874 microsomal glutathione S-transferase 2 /FL=gb:U77604.1 gb:Nm\_002413.1

205120\_s\_at gb:U29586.1 /DEF=Human beta-sarcoglycan dystrophin-associated glycoprotein mRNA, complete cds. /FEA=mRNA /PROD=dystrophin-associated glycoprotein /DB\_XREF=gi:1794188 /UG=Hs.77501 sarcoglycan, beta (43kD dystrophin-associated glycoprotein) /FL=gb:U31116.1 gb:U29586.1 gb:Nm\_000232.1

214880\_x\_at Consensus includes gb:D90453.1 /DEF=Human mRNA for I-caldesmon II. /FEA=mRNA /DB\_XREF=gi:219897 /UG=Hs.325474 caldesmon 1

205804\_s\_at gb:NM\_025228.1 /DEF=Homo sapiens hypothetical protein dJ434O14.3 (DJ434O14.3), mRNA. /FEA=mRNA /GEN=DJ434O14.3 /PROD=hypothetical protein dJ434O14.3 /DB\_XREF=gi:13435126 /UG=Hs.261373 hypothetical protein dJ434O14.3 /FL=gb:Nm\_025228.1

214696\_at Consensus includes gb:AF070569.1 /DEF=Homo sapiens clone 24659 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:3387938 /UG=Hs.29206 Homo sapiens clone 24659 mRNA sequence

218488\_at gb:NM\_020365.1 /DEF=Homo sapiens eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD) (EIF2B3), mRNA. /FEA=mRNA /GEN=EIF2B3 /PROD=eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD) /DB\_XREF=gi:9966778 /UG=Hs.283627 eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD) /FL=gb:AF257077.1 gb:Nm\_020365.1



Affy Probe_ID	Sequence Description
200795_at	gb:NM_004684.1 /DEF=Homo sapiens SPARC-like 1 (mast9, hev1n) (SPARCL1); mRNA. /FEA=mRNA /GEN=SPARCL1 /PROD=SPARC-like 1 /DB_XREF=gi:4758521 /UG=Hs.75445 SPARC-like 1 (mast9, hev1n) /FL=gb:NM_004684.1
206245_s_at	gb:NM_006469.1 /DEF=Homo sapiens NS1-binding protein (NS1-BP), mRNA. /FEA=mRNA /GEN=NS1-BP /PROD=NS1-binding protein /DB_XREF=gi:5453803 /UG=Hs.197298 NS1-binding protein /FL=gb:NM_006469.1
201297_s_at	Consensus includes gb:AK023321.1 /DEF=Homo sapiens cDNA.FLJ13259 fis, clone OVARC1000876, moderately similar to MOB1 PROTEIN. /FEA=mRNA /DB_XREF=gi:10435206 /UG=Hs.196437 hypothetical protein FLJ10788 /FL=gb:AB016839.1 gb:BC003398.1 gb:NM_018221.1
207168_s_at	gb:NM_004893.1 /DEF=Homo sapiens H2A histone family, member Y (H2AFY), mRNA. /FEA=mRNA /GEN=H2AFY /PROD=H2A histone family, member Y /DB_XREF=gi:4758495 /UG=Hs.75258 H2A histone family, member Y /FL=gb:AF054174.1 gb:NM_004893.1
57163_at	Cluster Incl. H93026:yv06a08.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-241910 /clone_end=3 /gb=H93026 /gi=1099354 /ug=Hs.25597 /len=600
201247_at	Consensus includes gb:BE513151 /FEA=EST /DB_XREF=gi:9720362 /DB_XREF=est:601171940F1 /CLONE=IMAGE:3545697 /UG=Hs.108689 sterol regulatory element binding transcription factor 2 /FL=gb:U02031.1 gb:NM_004599.1
207545_s_at	gb:NM_003744.1 /DEF=Homo sapiens numb (Drosophila) homolog (NUMB), mRNA. /FEA=mRNA /GEN=NUMB /PROD=numb (Drosophila) homolog /DB_XREF=gi:4505478 /UG=Hs.78890 numb (Drosophila) homolog /FL=gb:NM_003744.1 gb:L40393.1
208638_at	Consensus includes gb:BE910010 /FEA=EST /DB_XREF=gi:10406176 /DB_XREF=est:601498050F1 /CLONE=IMAGE:3899869 /UG=Hs.182429 protein disulfide isomerase-related protein /FL=gb:BC001312.1
203676_at	gb:NM_002076.1 /DEF=Homo sapiens glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID) (GNS), mRNA. /FEA=mRNA /GEN=GNS /PROD=glucosamine (N-acetyl)-6-sulfatase precursor /DB_XREF=gi:4504060 /UG=Hs.321070 glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID) /FL=gb:NM_002076.1
213283_s_at	Consensus includes gb:BG285616 /FEA=EST /DB_XREF=gi:13037752 /DB_XREF=est:602380622F1 /CLONE=IMAGE:4498325 /UG=Hs.79971 sal (Drosophila)-like 2.
210747_at	gb:M24364.1 /DEF=Human MHC class II lymphocyte antigen DQB mRNA, complete cds, haplotype DR7, DQw9. /FEA=mRNA /PROD=lymphocyte antigen /DB_XREF=gi:529041 /UG=Hs.73931 major histocompatibility complex, class II, DQ beta 1 /FL=gb:M24364.1

# Affy Probe\_ID

## Sequence Description

201062\_at gb:M81635.1 /DEF=Homo sapiens erythrocyte membrane protein mRNA, complete cds. /FEA=mRNA /GEN=stomatin peptide /PROD=stomatin peptide /DB\_XREF=gi:181183 /UG=Hs.160483 erythrocyte membrane protein band 7.2 (stomatin) /FL=gb:M81635.1 gb:NM\_004099.1

222315\_at Consensus includes gb:AW972855 /FEA=EST /DB\_XREF=gi:8162701 /DB\_XREF=est:EST384950 /UG=Hs.292853 ESTs

208511\_at gb:NM\_021000.1 /DEF=Homo sapiens pituitary tumor-transforming 3 (PTTG3), mRNA. /FEA=CDS /GEN=PTTG3 /PROD=pituitary tumor-transforming 3 /DB\_XREF=gi:10337606 /UG=Hs.247762 pituitary tumor-transforming 3 /FL=gb:NM\_021000.1 gb:AF095289.1

220644\_at gb:NM\_014137.1 /DEF=Homo sapiens PRO0650 protein (PRO0650), mRNA. /FEA=mRNA /GEN=PRO0650 /PROD=PRO0650 protein /DB\_XREF=gi:7662581 /UG=Hs.177258 PRO0650 protein /FL=gb:AF090941.1 gb:NM\_014137.1

36830\_at Cluster Incl. U80034:Human mitochondrial intermediate peptidase precursor (MIPEP) mRNA, mitochondrial gene encoding mitochondrial protein, complete cds /cds=(74,2215) /gb=U80034 /gi=1763641 /ug=Hs.68583 /len=2374

208286\_x\_at gb:NM\_002701.1 /DEF=Homo sapiens POU domain, class 5, transcription factor 1 (POU5F1), mRNA. /FEA=mRNA /GEN=POU5F1 /PROD=POU domain, class 5, transcription factor 1 /DB\_XREF=gi:4505966 /UG=Hs.2860 POU domain, class 5, transcription factor 1 /FL=gb:NM\_002701.1

222135\_at Consensus includes gb:AK022663.1 /DEF=Homo sapiens cDNA FLJ12601 fis, clone NT2RM4001414. /FEA=mRNA /DB\_XREF=gi:10434187 /UG=Hs.288549 Homo sapiens cDNA FLJ12601 fis, clone NT2RM4001414

218264\_at gb:NM\_016567.1 /DEF=Homo sapiens cdk inhibitor p21 binding protein (TOK-1), mRNA. /FEA=mRNA /GEN=TOK-1 /PROD=cdk inhibitor p21 binding protein /DB\_XREF=gi:7706580 /UG=Hs.279862 cdk inhibitor p21 binding protein /FL=gb:AB040450.1 gb:NM\_016567.1

202401\_s\_at gb:NM\_003131.1 /DEF=Homo sapiens serum response factor (c-fos serum response element-binding transcription factor) (SRF), mRNA. /FEA=mRNA /GEN=SRF /PROD=serum response factor (c-fos serum response element-binding transcription factor) /DB\_XREF=gi:4507204 /UG=Hs.155321 serum response factor (c-fos serum response element-binding transcription factor) /FL=gb:J03161.1 gb:NM\_003131.1

220322\_at gb:NM\_019618.1 /DEF=Homo sapiens interleukin-1 homolog 1 (IL-1H1), mRNA. /FEA=mRNA /GEN=IL-1H1 /PROD=interleukin-1 homolog 1 /DB\_XREF=gi:9665233 /UG=Hs.211238 interleukin-1 homolog 1 /FL=gb:AF206696.1 gb:AF200492.1 gb:NM\_019618.1

Affy Probe_ID	Sequence Description
202083_s_at	Consensus includes gb:AI017770 /FEA=EST /DB_XREF=gi:3232106 /DB_XREF=est:ov32b12.x1 /CLONE=IMAGE:1639007 /UG=Hs.75232 SEC14 (S. cerevisiae)-like 1 /FL=gb:D67029.1 gb:Nm_003003.1 gb:Nm_023926.1 /DEF=Homo sapiens hypothetical protein FLJ12895 (FLJ12895), mRNA. /FEA=mRNA /GEN=FLJ12895 /PROD=hypothetical protein FLJ12895 /DB_XREF=gi:12965194 /UG=Hs.235390 hypothetical protein FLJ12895 /FL=gb:Nm_023926.1
212151_at	Consensus includes gb:BF967998 /FEA=EST /DB_XREF=gi:12335213 /DB_XREF=est:602269506F1 /CLONE=IMAGE:435777 /UG=Hs.21851 Homo sapiens cDNA FLJ12900 fis, clone NT2RP2004321 gb:AF164794.1 /DEF=Homo sapiens Diff33 protein homolog mRNA, complete cds. /FEA=mRNA /PROD=Diff33 protein homolog /DB_XREF=gi:8895090 /UG=Hs.146668 KIAA1253 protein /FL=gb:AF164794.1
210879_s_at	gb:AF334812.1 /DEF=Homo sapiens Rab11 interacting protein Rip11a mRNA, complete cds. /FEA=mRNA /PROD=Rab11 interacting protein Rip11a /DB_XREF=gi:13377896 /UG=Hs.24557 KIAA0857 protein /FL=gb:AF334812.1
219252_s_at	gb:Nm_017856.1 /DEF=Homo sapiens hypothetical protein FLJ20514 (FLJ20514), mRNA. /FEA=mRNA /GEN=FLJ20514 /PROD=hypothetical protein FLJ20514 /DB_XREF=gi:8923480 /UG=Hs.44423 hypothetical protein FLJ20514 /FL=gb:Nm_017856.1
202599_s_at	gb:Nm_003489.1 /DEF=Homo sapiens nuclear receptor interacting protein 1 (NRIP1), mRNA. /FEA=mRNA /GEN=NRIP1 /PROD=receptor interacting protein 140 /DB_XREF=gi:4505454 /UG=Hs.155017 nuclear receptor interacting protein 1 /FL=gb:Nm_003489.1
213820_s_at	Consensus includes gb:T54159 /FEA=EST /DB_XREF=gi:656020 /DB_XREF=est:ya90f07.s2 /CLONE=IMAGE:68965 /UG=Hs.172803 hypothetical protein MGC10327
217650_x_at	Consensus includes gb:AI088162 /FEA=EST /DB_XREF=gi:3427138 /DB_XREF=est:oz96b09.x1 /CLONE=IMAGE:1683161 /UG=Hs.262292 ESTs, Moderately similar to ALU3_HUMAN ALU SUBFAMILY SB1 SEQUENCE CONTAMINATION WARNING ENTRY H.sapiens
222060_at	Consensus includes gb:AI357616 /FEA=EST /DB_XREF=gi:4109237 /DB_XREF=est:qu20g08.x1 /CLONE=IMAGE:1965374 /UG=Hs.101651 Homo sapiens mRNA; cDNA DKFZp434C107 (from clone DKFZp434C107)
204865_at	gb:Nm_005181.2 /DEF=Homo sapiens carbonic anhydrase III, muscle specific (CA3), mRNA. /FEA=mRNA /GEN=CA3 /PROD=carbonic anhydrase III /DB_XREF=gi:6996001 /UG=Hs.82129 carbonic anhydrase III, muscle specific /FL=gb:BC004897.1 gb:Nm_005181.2

Affy Probe_ID	Sequence Description
206739_at	gb:NM_018953.1 /DEF=Homo sapiens homeo box C5 (HOXC5), mRNA. /FEA=mRNA /GEN=HOXC5 /PROD=homeo box C5 /DB_XREF=gi:11321637 /UG=Hs.111473 homeo box C5 /FL=gb:NM_018953.1
203888_at	gb:NM_000361.1 /DEF=Homo sapiens thrombomodulin (THBD), mRNA. /FEA=mRNA /GEN=THBD /PROD=thrombomodulin /DB_XREF=gi:4507482 /UG=Hs.2030 thrombomodulin /FL=gb:M16552.1 gb:NM_000361.1
203367_at	gb:NM_007026.1 /DEF=Homo sapiens MKP-1 like protein tyrosine phosphatase (MKP-L), mRNA. /FEA=mRNA /GEN=MKP-L /PROD=MKP-1 like protein tyrosine phosphatase /DB_XREF=gi:5902001 /UG=Hs.91448 MKP-1 like protein tyrosine phosphatase /FL=gb:BC000370.1 gb:BC001894.1 gb:BC004448.1 gb:AF038844.1 gb:NM_007026.1 gb:AF120032.1
200615_s_at	Consensus includes gb:AL567295 /FEA=EST /DB_XREF=gi:12920509 /DB_XREF=est:AL567295 /CLONE=CS0DF027YO06 (3 prime) /UG=Hs.74626 adaptor-related protein complex 2, beta 1 subunit /FL=gb:M34175.1 gb:NM_001282.1
219736_at	gb:NM_018700.1 /DEF=Homo sapiens zinc-binding protein Rbcc728 (Rbcc728), mRNA. /FEA=mRNA /GEN=Rbcc728 /PROD=zinc-binding protein Rbcc728 /DB_XREF=gi:8924237 /UG=Hs.121429 zinc-binding protein Rbcc728 /FL=gb:NM_018700.1
213978_at	Consensus includes gb:AA338945 /FEA=EST /DB_XREF=gi:1991203 /DB_XREF=est:EST43994 /UG=Hs.319825
219132_at	Homo sapiens, clone IMAGE:3616574, mRNA, partial cds gb:NM_021255.1 /DEF=Homo sapiens pellino (Drosophila) homolog 2 (PELI2), mRNA. /FEA=mRNA /GEN=PELI2 /PROD=pellino (Drosophila) homolog 2 /DB_XREF=gi:10864062 /UG=Hs.44038 pellino (Drosophila) homolog 2 /FL=gb:AF302502.1 gb:NM_021255.1
217873_at	gb:NM_016289.1 /DEF=Homo sapiens MO25 protein (LOC51719), mRNA. /FEA=mRNA /GEN=LOC51719 /PROD=MO25 protein /DB_XREF=gi:7706480 /UG=Hs.6406 MO25 protein /FL=gb:AF113536.1 gb:NM_016289.1
37549_g_at	Cluster Incl. U87408:Human clone IMAGE-74593 unknown protein mRNA, partial cds /cds=(0,1362) /gb=U87408 /gi=1842104 /ug=Hs.79340 /len=1982
218476_at	gb:NM_007171.1 /DEF=Homo sapiens protein-O-mannosyltransferase 1 (POMT1), mRNA. /FEA=mRNA /GEN=POMT1 /PROD=protein-O-mannosyltransferase 1 /DB_XREF=gi:6005839 /UG=Hs.99654 protein-O-mannosyltransferase 1 /FL=gb:AF095136.1 gb:NM_007171.1

Affy Probe ID	Sequence Description
203117_s_at	gb:NM_014871.1 /DEF=Homo sapiens KIAA0710 gene product (KIAA0710), mRNA. /FEA=mRNA /GEN=KIAA0710 /PROD=KIAA0710 gene product /DB_XREF=gi:7662257 /UG=Hs.273397 KIAA0710 gene product /FL=gb:AB014610.1 gb:NM_014871.1
202655_at	gb:NM_006010.1 /DEF=Homo sapiens arginine-rich, mutated in early stage tumors (ARMET), mRNA. /FEA=mRNA /GEN=ARMET /PROD=arginine-rich protein /DB_XREF=gi:5174392 /UG=Hs.75412 arginine-rich, mutated in early stage tumors /FL=gb:M83751.1 gb:NM_006010.1
203304_at	gb:NM_012342.1 /DEF=Homo sapiens putative transmembrane protein (NMA), mRNA. /FEA=mRNA /GEN=NMA /PROD=putative transmembrane protein /DB_XREF=gi:6912533 /UG=Hs.78776 putative transmembrane protein /FL=gb:U23070.1 gb:NM_012342.1
213624_at	Consensus includes gb:AA873600 /FEA=EST /DB_XREF=gi:2969722 /DB_XREF=est:ob12a02.s1 /CLONE=IMAGE:1323434 /UG=Hs.42945 acid sphingomyelinase-like phosphodiesterase
220544_at	gb:NM_021733.1 /DEF=Homo sapiens testis-specific kinase substrate (TSKS), mRNA. /FEA=mRNA /GEN=TSKS /PROD=testis-specific kinase substrate /DB_XREF=gi:1119429 /UG=Hs.224210 testis-specific kinase substrate /FL=gb:NM_021733.1
203035_s_at	gb:NM_006099.1 /DEF=Homo sapiens protein inhibitor of activated STAT3 (PIAS3), mRNA. /FEA=mRNA /GEN=PIAS3 /PROD=protein inhibitor of activated STAT3 /DB_XREF=gi:5174628 /UG=Hs.76578 protein inhibitor of activated STAT3 /FL=gb:BC001154.1 gb:AB021868.1 gb:NM_006099.1
219055_at	gb:NM_018079.1 /DEF=Homo sapiens hypothetical protein FLJ10379 (FLJ10379), mRNA. /FEA=mRNA /GEN=FLJ10379 /PROD=hypothetical protein FLJ10379 /DB_XREF=gi:8922391 /UG=Hs.14229 hypothetical protein FLJ10379 /FL=gb:NM_018079.1
218497_s_at	gb:NM_002936.1 /DEF=Homo sapiens ribonuclease H1 (RNASEH1), mRNA. /FEA=mRNA /GEN=RNASEH1 /PROD=ribonuclease H1 /DB_XREF=gi:4506554 /UG=Hs.178655 ribonuclease H1 /FL=gb:BC002973.1
219635_at	gb:AF048994.1 gb:AF048995.1 gb:AF039652.1 gb:NM_002936.1
219635_at	gb:NM_025027.1 /DEF=Homo sapiens hypothetical protein FLJ14260 (FLJ14260), mRNA. /FEA=mRNA /GEN=FLJ14260 /PROD=hypothetical protein FLJ14260 /DB_XREF=gi:13430885 /UG=Hs.287629 hypothetical protein FLJ14260 /FL=gb:NM_025027.1
218689_at	gb:NM_022725.1 /DEF=Homo sapiens Fanconi anemia, complementation group F (FANCF), mRNA. /FEA=mRNA /GEN=FANCF /PROD=Fanconi anemia, complementation group F /DB_XREF=gi:12232376 /UG=Hs.65328 Fanconi anemia, complementation group F /FL=gb:NM_022725.1 gb:AF181994.1 gb:AF181995.1

**Affy Probe\_ID**

**Sequence Description**

205405\_at gb:NM\_003966.1 /DEF=Homo sapiens sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A (SEMA5A), mRNA. /FEA=mRNA /GEN=SEMA5A /PROD=sema domain, seven thrombospondin repeats (type1 and type 1-like), transmembrane domain (TM) and shortcytoplasmic domain, (semaphorin) 5A /DB\_XREF=gi:4506880 /UG=Hs.27621 sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A /FL=gb:U52840.1 gb:NM\_003966.1

213060\_s\_at Consensus includes gb:U58515.1 /DEF=Human chitinase (HUMTCHIT) mRNA, exon 1b form, partial cds. /FEA=mRNA /GEN=HUMTCHIT /PROD=chitinase /DB\_XREF=gi:1439567 /UG=Hs.154138 chitinase 3-like 2

212256\_at Consensus includes gb:BE906572 /FEA=EST /DB\_XREF=gi:10399901 /DB\_XREF=est:601502550F1 /CLONE=IMAGE:3904092 /UG=Hs.107260 hypothetical protein DKFZp586H0623

202860\_at gb:NM\_014856.1 /DEF=Homo sapiens KIAA0476 gene product (KIAA0476), mRNA. /FEA=mRNA /GEN=KIAA0476 /PROD=KIAA0476 gene product /DB\_XREF=gi:7662151 /UG=Hs.6684 KIAA0476 gene product /FL=gb:AB007945.1 gb:NM\_014856.1

209550\_at gb:U35139.1 /DEF=Human NECNIN related protein mRNA, complete cds. /FEA=mRNA /PROD=NECNIN related protein /DB\_XREF=gi:1754970 /UG=Hs.50130 necdin (mouse) homolog /FL=gb:NM\_002487.2 gb:U35139.1

202905\_x\_at Consensus includes gb:AI796269 /FEA=EST /DB\_XREF=gi:5361732 /DB\_XREF=est:wh44g10.x1 /CLONE=IMAGE:2383650 /UG=Hs.25812 Nijmegen breakage syndrome 1 (nibrin) /FL=gb:AF058696.1 gb:AF051334.1 gb:NM\_002485.2

217528\_at Consensus includes gb:BF003134 /FEA=EST /DB\_XREF=gi:10703409 /DB\_XREF=est:7g55f09.x1 /CLONE=IMAGE:3310409 /UG=Hs.72365 ESTs

206512\_at gb:NM\_005083.2 /DEF=Homo sapiens U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 1 (U2AF1RS1), mRNA. /FEA=mRNA /GEN=U2AF1RS1 /PROD=U2 small nuclear ribonucleoprotein auxiliaryfactor, small subunit-related protein 1 /DB\_XREF=gi:13186299 /UG=Hs.103962 U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 1 /FL=gb:D49676.1 gb:NM\_005083.2

211813\_x\_at gb:AF138303.1 /DEF=Homo sapiens decorin D mRNA, complete cds, alternatively spliced. /FEA=CDS /PROD=decorin D /DB\_XREF=gi:7381210 /UG=Hs.76152 decorin /FL=gb:AF138303.1

201530\_x\_at gb:NM\_001416.1 /DEF=Homo sapiens eukaryotic translation initiation factor 4A, isoform 1 (EIF4A1), mRNA. /FEA=mRNA /GEN=EIF4A1 /PROD=eukaryotic translation initiation factor 4A,isoform 1 /DB\_XREF=gi:4503528 /UG=Hs.129673 eukaryotic translation initiation factor 4A, isoform 1 /FL=gb:NM\_001416.1

# Affy Probe\_ID

## Sequence Description

209396\_s\_at gb:M80927.1 /DEF=Human glycoprotein mRNA, complete cds. /FEA=mRNA /PROD=glycoprotein /DB\_XREF=gi:348911 /UG=Hs.75184 chitinase 3-like 1 (cartilage glycoprotein-39) /FL=gb:M80927.1 gb:NM\_001276.1

218358\_at gb:NM\_024324.1 /DEF=Homo sapiens hypothetical protein MGC11256 (MGC11256), mRNA. /FEA=mRNA /GEN=MGC11256 /PROD=hypothetical protein MGC11256 /DB\_XREF=gi:13236568 /UG=Hs.28029 hypothetical protein MGC11256 /FL=gb:BC002894.1 gb:NM\_024324.1

204719\_at gb:NM\_007168.1 /DEF=Homo sapiens ATP-binding cassette, sub-family A (ABC1), member 8 (ABCA8), mRNA. /FEA=mRNA /GEN=ABCA8 /PROD=ATP-binding cassette, sub-family A member 8 /DB\_XREF=gi:6005700 /UG=Hs.38095 ATP-binding cassette, sub-family A (ABC1), member 8 /FL=gb:AB020629.1 gb:NM\_007168.1

222149\_x\_at Consensus includes gb:AL137398.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K052 (from clone DKFZp434K052). /FEA=mRNA /DB\_XREF=gi:6807944 /UG=Hs.169639 Homo sapiens mRNA; cDNA DKFZp434K052 (from clone DKFZp434K052)

214658\_at Consensus includes gb:BG286537 /FEA=EST /DB\_XREF=gi:13039504 /DB\_XREF=est:602382536F1 /CLONE=IMAGE:4500129 /UG=Hs.278391 CGI-109 protein

201212\_at gb:D55696.1 /DEF=Homo sapiens mRNA for cysteine protease, complete cds. /FEA=mRNA /PROD=cysteine protease /DB\_XREF=gi:1890049 /UG=Hs.18069 protease, cysteine, 1 (legumain) /FL=gb:BC003061.1 gb:D55696.1 gb:NM\_005606.1

208336\_s\_at gb:NM\_004868.1 /DEF=Homo sapiens glycoprotein, synaptic 2 (GPSN2), mRNA. /FEA=mRNA /GEN=GPSN2 /PROD=glycoprotein, synaptic 2 /DB\_XREF=gi:4759061 /UG=Hs.306122 glycoprotein, synaptic 2 /FL=gb:AF038959.1 gb:NM\_004868.1

217551\_at Consensus includes gb:AA719797 /FEA=EST /DB\_XREF=gi:2732896 /DB\_XREF=est:zg54c07.s1 /CLONE=IMAGE:397164 /UG=Hs.194362 olfactory receptor, family 7, subfamily E, member 13 pseudogene

203410\_at gb:NM\_006803.1 /DEF=Homo sapiens adaptor-related protein complex 3, mu 2 subunit (AP3M2), mRNA. /FEA=mRNA /GEN=AP3M2 /PROD=adaptor-related protein complex 3; mu 2 subunit /DB\_XREF=gi:5802999 /UG=Hs.77770 adaptor-related protein complex 3, mu 2 subunit /FL=gb:NM\_006803.1 gb:D38293.1

207621\_s\_at gb:NM\_007169.1 /DEF=Homo sapiens phosphatidylethanolamine N-methyltransferase (PEMT), mRNA. /FEA=mRNA /GEN=PEMT /PROD=phosphatidylethanolamine N-methyltransferase /DB\_XREF=gi:6005827 /UG=Hs.15192 phosphatidylethanolamine N-methyltransferase /FL=gb:AB029821.1 gb:AF176806.1 gb:NM\_007169.1

**Affy Probe ID**

**Sequence Description**

203588\_s\_at

Consensus includes gb:BG034328 /FEA=EST /DB\_XREF=gi:12427528 /DB\_XREF=est:602302213F1 /CLONE=IMAGE:4403787 /UG=Hs.19131 transcription factor Dp-2 (E2F dimerization partner 2) /FL=gb:NM\_006286.1 gb:U18422.1

221567\_at

gb:AF064599.1 /DEF=Homo sapiens nucleolar protein Nop30 (NOP) mRNA, alternatively spliced, complete cds. /FEA=mRNA /GEN=NOP /PROD=nucleolar protein Nop30 /DB\_XREF=gi:3172418 /UG=Hs.278439 nucleolar protein 3 (apoptosis repressor with CARD domain) /FL=gb:AF043244.1 gb:AF064599.1 gb:AF064600.1 gb:NM\_003946.1 gb:U90304.1 /DEF=Human iroquois-class homeodomain protein IRX-2a mRNA, complete cds. /FEA=mRNA /PROD=iroquois-class homeodomain protein IRX-2a /DB\_XREF=gi:1899219 /UG=Hs.25351 iroquois homeobox protein 5 /FL=gb:U90304.1 gb:NM\_005853.1

210239\_at

200797\_s\_at

Consensus includes gb:AI275690 /FEA=EST /DB\_XREF=gi:3897964 /DB\_XREF=est:qw03a03.x1 /CLONE=IMAGE:1989964 /UG=Hs.86386 myeloid cell leukemia sequence 1 (BCL2-related) /FL=gb:NM\_021960.1 gb:AF118124.1

202561\_at

Consensus includes gb:AF070613.1 /DEF=Homo sapiens clone 24585 mRNA sequence. /FEA=mRNA /DB\_XREF=gi:3387995 /UG=Hs.131814 tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase /FL=gb:AF082556.1 gb:NM\_003747.1

206502\_s\_at

gb:NM\_002196.1 /DEF=Homo sapiens insulinoma-associated 1 (INSM1), mRNA. /FEA=mRNA /GEN=INSM1 /PROD=insulinoma-associated 1 /DB\_XREF=gi:4504712 /UG=Hs.89584 insulinoma-associated 1 /FL=gb:M93119.1 gb:NM\_002196.1

206188\_at

gb:NM\_014789.1 /DEF=Homo sapiens KIAA0628 gene product (KIAA0628), mRNA. /FEA=mRNA /GEN=KIAA0628 /PROD=KIAA0628 gene product /DB\_XREF=gi:7662213 /UG=Hs.43133 KIAA0628 gene product /FL=gb:AB014528.1 gb:NM\_014789.1

217122\_s\_at

Consensus includes gb:AL031282 /DEF=Human DNA sequence from clone 283E3 on chromosome 1p36.21-36.33. Contains the alternatively spliced gene for Matrix Metalloproteinase in the Female Reproductive tract MIFR1, -2, MMP2122A, -B and -C, a novel gene, the alternatively spliced CDC2L2 ... /FEA=mRNA\_6 /DB\_XREF=gi:3860395 /UG=Hs.214646 KIAA0447 gene product

219432\_at

gb:NM\_014556.1 /DEF=Homo sapiens Ellis van Creveld syndrome (EVC), mRNA. /FEA=mRNA /GEN=EVC /PROD=Ellis van Creveld syndrome protein /DB\_XREF=gi:7657072 /UG=Hs.274446 Ellis van Creveld syndrome /FL=gb:AF216184.1 gb:AF239742.1 gb:NM\_014556.1



**Affy Probe ID**

**Sequence Description**

207266\_x\_at gb:NM\_016837.1 /DEF=Homo sapiens RNA binding motif, single stranded interacting protein 1 (RBMS1), transcript variant M5SP-3, mRNA. /FEA=mRNA /GEN=RBMS1 /PROD=RNA binding motif, single stranded interacting protein 1, isoform e /DB\_XREF=gi:8400719 /UG=Hs.241567 RNA binding motif, single stranded interacting protein 1 /FL=gb:NM\_016837.1

209676\_at gb:J03225.1 /DEF=Human lipoprotein-associated coagulation inhibitor mRNA, complete cds. /FEA=mRNA /GEN=TFPI /DB\_XREF=gi:180545 /UG=Hs.170279 tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) /FL=gb:J03225.1 gb:NM\_006287.2

209055\_s\_at Consensus includes gb:AW268817 /FEA=EST /DB\_XREF=gi:6655847 /DB\_XREF=est:xv38c01.x1 /CLONE=IMAGE:2815392 /UG=Hs.155174 CDC5 (cell division cycle 5, S. pombe, homolog)-like /FL=gb:NM\_001253.1 gb:U86753.1 gb:AB007892.1

206874\_s\_at Consensus includes gb:AL138761 /DEF=Human DNA sequence from clone RP11-16H23 on chromosome 10. Contains the gene KIAA0204 (HSLK) for a protein kinase, the COL17A1 gene for collagen type XVII alpha 1 (BP180), ESTs and GSSs /FEA=mRNA\_2 /DB\_XREF=gi:8573811 /UG=Hs.105751 Ste20-related serine/threonine kinase /FL=gb:D86959.1 gb:NM\_014720.1

221139\_s\_at gb:NM\_015989.1 /DEF=Homo sapiens cysteine sulfinic acid decarboxylase-related protein 2 (CSAD), mRNA. /FEA=mRNA /GEN=CSAD /PROD=cysteine sulfinic acid decarboxylase-related protein 2 /DB\_XREF=gi:7705333 /UG=Hs.279815 cysteine sulfinic acid decarboxylase-related protein 2 /FL=gb:AF116546.1 gb:NM\_015989.1

204136\_at gb:NM\_000094.1 /DEF=Homo sapiens collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive) (COL7A1), mRNA. /FEA=mRNA /GEN=COL7A1 /PROD=collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive) /DB\_XREF=gi:4502960 /UG=Hs.1640 collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive) /FL=gb:NM\_000094.1 gb:L02870.1

217841\_s\_at gb:NM\_016147.1 /DEF=Homo sapiens protein phosphatase methyltransferase-1 (PME-1), mRNA. /FEA=mRNA /GEN=PME-1 /PROD=protein phosphatase methyltransferase-1 /DB\_XREF=gi:7706644 /UG=Hs.63304 protein phosphatase methyltransferase-1 /FL=gb:BC003046.1 gb:AF157028.1 gb:NM\_016147.1

202735\_at gb:NM\_006579.1 /DEF=Homo sapiens emopamil-binding protein (sterol isomerase) (EBP), mRNA. /FEA=mRNA /GEN=EBP /PROD=emopamil-binding protein (sterol isomerase) /DB\_XREF=gi:5729809 /UG=Hs.75105 emopamil-binding protein (sterol isomerase) /FL=gb:NM\_006579.1

Affy Probe_ID	Sequence Description
200625_s_at	gb:NM_006367.2 /DEF=Homo sapiens adenylyl cyclase-associated protein (CAP), mRNA. /FEA=mRNA /GEN=CAP /PROD=adenylyl cyclase-associated protein /DB_XREF=gi:10938021 /UG=Hs.104125 adenylyl cyclase-associated protein /FL=gb:NM_006367.2 gb:L12168.1 gb:M98474.1
204288_s_at	gb:NM_021069.1 /DEF=Homo sapiens ArgAbl-interacting protein ArgBP2 (ARGBP2), transcript variant 2, mRNA. /FEA=mRNA /GEN=ARGBP2 /PROD=ArgAbl-interacting protein 2, isoform 2 /DB_XREF=gi:10947117 /UG=Hs.278626 ArgAbl-interacting protein ArgBP2 /FL=gb:NM_021069.1 gb:AB018320.1
211626_x_at	gb:M21535.1 /DEF=Human erg protein (ets-related gene) mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:182182 /FL=gb:M21535.1
209605_at	gb:D87292.1 /DEF=Homo sapiens mRNA for rhodanese, complete cds. /FEA=mRNA /PROD=rhodanese /DB_XREF=gi:1877030 /UG=Hs.248267 thiosulfate sulfotransferase (rhodanese) /FL=gb:D87292.1
212560_at	Consensus includes gb:AV728268 /FEA=EST /DB_XREF=gi:10837689 /DB_XREF=est:AV728268 /CLONE=HTCACH05 /UG=Hs.82845 Homo sapiens cDNA: FLJ21930 fis, clone HEP04301, highly similar to HSU90916 Human clone 23815 mRNA sequence